

1 **The CaMV 35S promoter is the most widely used promoter in plant biotechnology - despite**  
2 **being derived from a pathogenic virus. How and why did that happen? Here's...**

## 3 **A Short History of the CaMV 35S Promoter**

4 **Marc Somssich**

5 Person Lab, School of BioSciences, the University of Melbourne, Parkville 3010, VIC, Australia

6 Email: [marc.somssich@unimelb.edu.au](mailto:marc.somssich@unimelb.edu.au) ; Twitter: [@somssichm](https://twitter.com/somssichm)

7

### 8 **The Mosaic Disease (1921 – 1937)**

9 The effects of the Cauliflower mosaic virus (CaMV) were first noted in **1921** in Chinese cabbage,  
10 where it caused mosaic-like necrotic lesions on leaf surfaces<sup>1</sup>. In the following years, the disease  
11 was regularly observed on Chinese cabbage, turnip or pot-herb mustard<sup>2</sup>. However, it was only in  
12 the **1930s**, that scientists invested more time and resources to investigate the disease<sup>2</sup>. This was  
13 spawned by severe yield losses on cabbage fields in the American Midwest, which strikingly  
14 were preceded by a heavy infestation of the field with aphids<sup>2</sup>. During this time, similar mosaic-  
15 like lesions were found on cauliflower in California, prompting investigations if these were  
16 caused by the same disease<sup>3</sup>. In a **1937** study, using infected cauliflower plants collected in  
17 California, C. M. Tompkins found that he could transmit the disease from the infected  
18 cauliflower plants to 51 different vegetable varieties, all belonging to the crucifer family  
19 (Cruciferae/Brassicaceae)<sup>3</sup>. This family includes, e.g., cauliflower, broccoli, cabbage, kale,  
20 turnip, kohlrabi or Chinese cabbage<sup>3</sup>. In the same study, he found that at least three different  
21 aphids can function as insect vectors, all three common inhabitants of cauliflower crop fields,  
22 thereby indicating that it is indeed a virus that causes the disease<sup>3</sup>. Although originally referred to  
23 simply as the ‘mosaic disease’, the virus was named Cauliflower mosaic virus, due to its  
24 described isolation from cauliflower<sup>3</sup>.

### 25 **The Cauliflower Mosaic Virus (1937 – 1978)**

26 In the late **1940s**, research on the CaMV intensified once more, this time primarily in Europe,  
27 where it caused devastating losses in cauliflower and broccoli harvests across Great Britain<sup>4</sup>. As  
28 this was just after the end of World War II the impact was especially dramatic, as food was

29 already a scarcity<sup>4</sup>. One of the first important findings in the following years was that CaMV is a  
30 non-circulative (and non-persistent) virus, meaning that it does not enter its aphid vector, but just  
31 ‘sticks’ to the insects stylet, and is thereby transported from an infected plant to a healthy one<sup>5</sup>.  
32 Interestingly, in 2007, researchers were able to pinpoint the exact position, at which the virus is  
33 perceived by the insect, an area roughly 5  $\mu\text{m}$  long and less than 1  $\mu\text{m}$  wide at the most distal tip  
34 of the aphid’s stylet<sup>6</sup>. Through another important finding in the **1960s**, the CaMV was identified  
35 as the first plant virus containing double-stranded DNA<sup>7</sup>. This is of particular importance,  
36 because this feature is a pre-requisite for the viral DNA to be transcribed in plant cells<sup>7</sup>.  
37 Furthermore, this was the first indication that CaMV is a pararetrovirus (in contrast to the more  
38 commonly known single-strand RNA-containing retroviruses), even though this was only  
39 determined much later<sup>8</sup>. In **1980**, the whole genome (8024 double-stranded, circular base pairs  
40 (bp)) of the virus was annotated and found to contain six putative open reading frames<sup>9,10</sup>. At this  
41 point, scientists started to focus on deciphering the molecular details of plant infection by the  
42 virus. In the early 1980s it was discovered that the six coding regions are transcribed as only two  
43 mRNAs, the short, monocistronic 19S RNA, and the whole-genome covering 35S mRNA<sup>11</sup>.  
44 While the 19S RNA encodes a single protein, which was later found to be involved in gene  
45 silencing suppression in the host cell, the long 35S RNA serves as a template for whole genome  
46 replication, and is furthermore spliced into four individual mRNAs<sup>12–14</sup>. The 35S RNA also has  
47 two very curious features; (I) although serving also as a template for the genome, it is actually  
48 longer than the genome, as the 5’ and 3’ ends overlap by 200 nucleotides (nt); and (II) it has an  
49 unusually long 600 nt leader sequence<sup>15</sup>. This 600 nt leader was later found to be transcribed into  
50 ‘massive amounts’ of 21 to 24 nt sense and antisense RNAs, which could function as ‘decoys’  
51 during infection, to divert the host cell’s silencing machinery from the actual coding 35S  
52 mRNAs<sup>16</sup>. However, the most important finding during that period was that the 19S and 35S  
53 reading frames were found to be highly expressed in infected plant cells, implying that the virus  
54 must have inserted its own double-stranded DNA into the plant cell, and that this inserted piece  
55 of viral DNA must contain all elements necessary to initiate transcription at high levels in host  
56 cells<sup>11,15</sup>.

### 57 **The Cauliflower Mosaic Virus meets Plant Biotechnology (1978 – 1985)**

58 At this point it is important to note that in the late 1970s and early **1980s**, the field of plant  
59 molecular biology and genetics/genomics was still in its infancy<sup>17</sup>. *Arabidopsis thaliana* had just

60 been accepted as a model system (see also ‘A Short History of *Arabidopsis thaliana* (L.) Heynh.  
61 Columbia-0’<sup>18</sup>)<sup>17</sup>. Direct transformation of a living plant with a transgene was not possible yet  
62 and very few individual genes had been cloned or studied at all. Furthermore, only a single  
63 promoter functional in plants had been fully described – the bacterial *octopine synthase* gene  
64 promoter, while a single promoter from plants, the pea *Ribulose-1,5-bisphosphate carboxylase*  
65 small subunit promoter, had been roughly located<sup>17,19,20</sup>. Accordingly, when it became clear that  
66 CaMV inserted its DNA into plant cells, and that this DNA was then expressed at high levels,  
67 plant biologists immediately recognized the potential use of CaMV as a cloning vector for plant  
68 transformation, and for expressing their genes of interest in the plant<sup>11,21,22</sup>. This resulted in two  
69 paths of research: First, researchers tried to insert a foreign gene into the genome of CaMV to  
70 determine whether this will get inserted and expressed in the host cell as well. Secondly, they  
71 attempted to identify the exact DNA sequences responsible for the strong expression of the  
72 CaMV genes in plant cells. Regarding the first path, researchers quickly progressed, and in the  
73 mid-eighties had successfully cloned bacterial and mammalian genes into the CaMV genome,  
74 and demonstrated that these genes were then transferred and expressed in plant cells<sup>23–25</sup>.  
75 However, they also realized that CaMV would only tolerate the insertion of short DNA fragments  
76 (~250 bp), and with the recent establishment of *Agrobacterium*-mediated plant transformation  
77 (see also ‘A Short History of Plant Transformation’<sup>26</sup>), the research in CaMV-mediated plant  
78 transformation was subsequently abandoned in the early 1990s<sup>27,28</sup>. However, the second research  
79 path, the identification of the exact sequences that control gene expression in CaMV-infected  
80 cells, turned out to be a much bigger success.

### 81 **The CaMV 35S Promoter (1985-2000)**

82 Up until **1985** it was almost impossible to over- or misexpress a gene of interest *in planta* –  
83 nowadays an invaluable and indispensable tool to study the function of a specific gene. The  
84 identification of the CaMV 35S promoter would finally change this. In order to define the exact  
85 sequences controlling viral gene expression *in planta*, researchers first created several deletion  
86 variants of the roughly 1000 bp promoter region of the 35S gene and fused these variants  
87 upstream of the *human growth hormone (hgh)* gene<sup>29</sup>. Notably, they used *Agrobacterium*  
88 *tumefaciens* to transform plant cells with their 35S::hgh variants, not the CaMV itself<sup>29</sup>. They  
89 found that DNA sequences 46 bp upstream of the 35S gene resulted in minimal expression, while  
90 a 343 bp fragment led to strong gene expression across all plant tissues tested<sup>29</sup>. The full 343 bp

91 segment was therefore designated the ‘CaMV 35S promoter’, while the 46 bp segment was  
92 considered as the so-called ‘minimal promoter’<sup>29</sup>. In follow-up studies, these 343 bp could then  
93 be further subdivided into several individual stretches, which would promote expression in  
94 different cell types or tissues, in either an additive or combinatorial fashion<sup>30,31</sup>. Based on these  
95 groundbreaking findings, numerous versions of the promoter emerged over the course of the  
96 following years; for example, simply placing two CaMV35S promoters in a tandem led to  
97 enhanced strength of the expression system<sup>32</sup>. Furthermore, the 46 bp minimal promoter also  
98 proved to be a highly useful tool: In the following years, short regulatory sequences within  
99 different gene promoters were identified that were bound by specific transcriptional activators<sup>33–</sup>  
100 <sup>36</sup>. By combining these activating elements with the minimal 35S promoter, scientists generated  
101 promoters that could promote gene expression in combination with the right activator<sup>33–36</sup>. This  
102 could be a plant hormone, such as auxin, therefore activating gene expression in a pattern  
103 reflecting the endogenous auxin concentration, or in response to external addition of the  
104 hormone<sup>33</sup>. Moreover, effectors could also be animal hormones such as estrogen and  
105 glucocorticoid, or ethanol, all of which are normally not present in the plant, therefore giving  
106 researchers complete control over when and where expression could be induced<sup>35–37</sup>. Finally, by  
107 combining the minimal 35S promoter with transcriptional binding sites for pathogen-responsive  
108 activators, researchers could engineer constructs that would confer enhanced resistance in the  
109 event of a pathogen attack<sup>34</sup>.

110 This last point was not the main finding that made the CaMV 35S promoter so appealing to crop  
111 scientists, however. In **1986**, merely a year after the CaMV 35S promoter was correctly  
112 described, it was used to promote expression of the *5-enolpyruvylshikimate-3-phosphate synthase*  
113 (*EPSP*) gene in transgenic petunia<sup>38</sup>. The EPSP is an essential enzyme in the aromatic amino acid  
114 biosynthetic pathway<sup>39</sup>. And this enzyme is also the specific target for the herbicide glyphosate<sup>40</sup>.  
115 Accordingly, plants that overexpress the *EPSP* gene from the 35S promoter acquire an increased  
116 tolerance towards glyphosate treatment<sup>38</sup>. This successful engineering of the first transgenic  
117 herbicide-tolerant plants combined two major scientific breakthroughs of the early 1980s – the  
118 establishment of *Agrobacterium*-mediated plant transformation and the identification of the  
119 CaMV 35S promoter – and together these three milestones, all published within three years,  
120 meant a giant leap forward for both the plant science community and the developing field of plant  
121 biotechnology<sup>29,38,41</sup>. Over the course of the following 20 to 30 years, the 35S promoter became

122 the most frequently used promoter in plant biotechnology, and almost every genetically modified  
123 crop plant that made it into our fields carries a version of this promoter<sup>42</sup>.

#### 124 **The CaMV 35S Promoter as Target for Anti-GE Activists (1990-today)**

125 The creation of genetically engineered crop plants not only gave a boost to plant science, it also  
126 activated the anti-GE (Genetic Engineering) movement. And in the 1990s, the 35S promoter  
127 became one of their main targets<sup>43</sup>. Interestingly though, there was no biosafety-incident or  
128 something comparable, that spawned a reasonable fear of the 35S promoter – it was a  
129 combination of insufficient outreach and bad public relations work from the scientific  
130 community, and the mere origin of the 35S promoter from a pathogenic virus<sup>42</sup>. There were,  
131 however, two incidents that clearly contributed to tarnishing the reputation of the 35S promoter in  
132 the public eye: the Petunia field trial in Germany, in **1990**, and the Pusztai affair in Great Britain,  
133 in 1998<sup>44,45</sup>.

134 In the 1980s, researchers at the Max-Planck Institute for Plant Breeding Research (MPIPZ) in  
135 Cologne, Germany, were working on transposable elements in maize (so called ‘jumping genes’  
136 or ‘transposons’)<sup>46,47</sup>. They found that white flowers from petunia could be converted into salmon  
137 red flowers, by introducing a maize transgene under control of the CaMV 35S promoter<sup>46</sup>. They  
138 then used these red flowering plants in a field trial to identify transposons in petunia, arguing that  
139 if enough plants are sowed out, the rare ‘jump’ of a transposon into the introduced maize  
140 transgene would be readily identified<sup>48</sup>. This insertion event should render the maize transgene  
141 inactive, thereby turning the red flower back to white – a clear visible sign that a transposon had  
142 been ‘trapped’<sup>48</sup>. The chance for this to happen was estimated to be around 0.0001 %, and so the  
143 expectation was that only a few individual flowers in the population of over 30 000 plants would  
144 revert back to their white color<sup>48</sup>. The result, however, was a reversion rate of almost 60 %<sup>48</sup>. The  
145 researchers later discovered that this was mostly due to epigenetic gene silencing, the auto-  
146 inactivation of gene expression - a protective mechanisms of the plant cell if expression appears  
147 to get out of control<sup>48,49</sup>. The plant cell had simply turned off the CaMV 35S promoter because it  
148 was too strong, which also demonstrates why the CaMV had evolved its 19S protein as a  
149 silencing suppressor - something only discovered many years later in 2007<sup>12,48</sup>. While this was an  
150 exciting but surprise finding for the plant science community (‘epigenetic gene silencing’ was not  
151 well studied or understood at that time), it also became a public relations problem for the

152 MPIPZ<sup>44</sup>. The petunia experiment was the first field trial with transgenic plants in Germany, a  
153 country that is notoriously critical and reserved when it comes to genetic engineering<sup>44,50,51</sup>. The  
154 trial therefore was accompanied by protests from anti-GE activists claiming that the scientists did  
155 not understand genetic engineering well enough to undertake such an experiment outside of the  
156 controlled environment of a green house without considerable risks<sup>51</sup>. The scientists, on the other  
157 side, were both interested in the scientific outcome of the experiment, but also to demonstrate to  
158 the public that they were able to control such an experiment<sup>44,51</sup>. Needless to say, the results did  
159 not go over well with the public, and the protesters felt reassured of their claim<sup>44</sup>. In the following  
160 two decades, there were some further field trials of GE-plants in Germany, but as of 2015, 75 %  
161 of Germans are still opposed to growing genetically modified crops and since 2013 no further  
162 field trials with GE crops were approved ([www.bvl.bund.de/EN/](http://www.bvl.bund.de/EN/))<sup>52</sup>. Interestingly, more than 25  
163 years after the petunia trial, in 2017, red-colored commercial petunias were recalled from stores  
164 worldwide when it was discovered that they were actually transgenic – they carried the same  
165 maize transgene that was used at the MPIPZ in 1987<sup>53,54</sup>. This episode was widely publicized in  
166 Germany at the time, and certainly contributed to a majority of Germans still remaining critical  
167 when it comes to genetic engineering<sup>52</sup>. However, even though the 35S promoter was indeed  
168 indirectly responsible for this public relations debacle for the German plant science community,  
169 at the time, the 35S promoter was not yet singled out as a threat to the environment and human  
170 health, and the activists focused on genetic engineering as a whole instead. This would change  
171 with the Pusztai affair in **1998**<sup>55</sup>.

172 In the wake of increasing consumer concerns over the safety of genetically modified food,  
173 renowned protein scientist Árpád Pusztai, who was in the midst of conducting the first major  
174 study on possible health effects of transgenic crops, was interviewed on British TV about his  
175 ongoing experiments<sup>56</sup>. He stated that the trials included rats that were fed genetically modified  
176 potatoes and that they seemed to be less healthy than rats that were fed the unmodified  
177 counterpart<sup>56</sup>. He also acknowledged that he could not tell what caused these effects, and that he  
178 actually had concerns about the experimental design and the controls included<sup>56</sup>. Accordingly, he  
179 stated that more testing was needed until any firm conclusions could be reached<sup>56</sup>. However, he  
180 then went on to state that, if given the choice to eat transgenic crops now he ‘*wouldn't eat it*’,  
181 adding that he thought it was ‘*very, very unfair to use our fellow citizens as guinea pigs*’<sup>56,57</sup>. Not  
182 surprisingly, this last sentence resulted in a major pushback against genetic engineering from the

183 public, which was further exacerbated by how the Rowett Research Institute (Pusztai's employer)  
184 subsequently handled the situation<sup>56,58</sup>. Overwhelmed by the massive backlash the institute  
185 received from the public and media, the Rowett Institute panicked and shut down Pusztai's lab,  
186 collected all lab books, suspended him indefinitely, and, worst of all, forbade him from talking to  
187 the press<sup>56,57</sup>. The director also released a statement, in which he described Pusztai's data as 'a  
188 total muddle', and apologized for releasing 'misleading information', before an investigation into  
189 the case had even begun<sup>58</sup>. This reaction drew massive criticism from both, fellow scientists and  
190 the public<sup>57</sup>. The scientific community, who valued Pusztai as a highly reputable colleague, and  
191 certainly an authority in the field, were shocked by his harsh treatment, and noted that '*it is an*  
192 *unacceptable code of practice by the Rowett and its Director, Professor James, to set themselves*  
193 *up as arbiters or judges of the validity of the data which could have such a profound importance*  
194 *not only for scientists, but also for the public and its health*'<sup>57</sup>. For members of the public,  
195 similarly, the actions taken by the institute seemed as if the Institute was trying to silence a  
196 dissident member, and cover up his findings<sup>57,58</sup>.

197 So what was all of this about? In his experiments, Pusztai fed transgenic potatoes expressing a  
198 snowdrop lectin (*Galanthus nivalis agglutinin (GNA)*) to rats, to test for any effects this would  
199 have on their health<sup>59</sup>. Plant lectins are sugar-binding proteins involved in cell immunity by  
200 detecting specific sugar chains on the surfaces of proteins, but also on viruses or bacteria<sup>60</sup>. GNA  
201 was shown to be toxic to some insects, among them several major crop pests<sup>61</sup>. So in this case,  
202 binding of GNA to the insect pest would kill the insect<sup>61</sup>. At the same time, GNA was shown not  
203 to be toxic to mammals, including rats (as demonstrated by Pusztai himself in 1990)<sup>62</sup>. Based on  
204 these findings the idea was developed to create transgenic crops expressing *GNA*, to enhance  
205 their tolerance to insect pests<sup>60</sup>. One of the first such crop plants was a potato expressing the *GNA*  
206 from the CaMV 35S promoter<sup>63</sup>. Pusztai's feeding trial was now intended to check for any health  
207 risks for mammals coming from such transgenic crop plants. In order to do this, he fed the rats  
208 either transgenic potatoes, non-transgenic potatoes, or non-transgenic potatoes that were laced  
209 with GNA<sup>59</sup>. And what he found was that several of the rats that were fed the transgenic potatoes,  
210 were less healthy than the rats that were fed the unmodified counterpart, or the unmodified but  
211 GNA-laced potatoes<sup>59</sup>. However, the observed effects were also highly variable from potato to  
212 potato, and most effects could not be traced back to the expressed GNA, as GNA-  
213 supplementation, even at high concentrations, did not result in the same effects on the rats as

214 *GNA* transgene expression<sup>57,59,64</sup>. Therefore, Pusztai speculated that most of the effects were not  
215 due to the expressed GNA protein, but to the transgene itself, the position where it was inserted,  
216 the transformation procedure, or general alterations in the composition of the potato caused by  
217 the procedure to obtain the genetically modified plant<sup>59,64</sup>. Eventually, further analysis of  
218 Pusztai's data by external experts and an investigation by a commission set up by the Royal  
219 British Society all found that the Pusztai experimental setup was indeed clearly flawed, and that  
220 for this reason, no conclusions could be drawn from his findings<sup>57,64-66</sup>. They showed, for  
221 example, that the nutritional value varied widely between the different transgenic potatoes, as  
222 well as to their parental line, which already makes it impossible to distinguish if any effects were  
223 caused by the transgene, or simply were due to this variation<sup>64</sup>. Furthermore, as this diet is not  
224 suitable for rats, the animals were all protein-starved, affecting their general physiology, again  
225 making it impossible to trace any effects back to the transgene<sup>64</sup>. Also, the controls were flawed.  
226 The transgenic potatoes were created by transformation of cultured cells and regeneration of  
227 plants from these cells<sup>63</sup>. However, the wild type control plants had not gone through such a  
228 procedure<sup>57,64,65</sup>. This is especially important, as plants derived from tissue culture exhibit  
229 somaclonal variation, which can account for a wide spectrum of effects, most of all the observed  
230 differences in nutritional value between the different potatoes<sup>57,64,65,67</sup>. And there are many more  
231 problems with the study as detailed in the various reports<sup>57,64-67</sup>. Thus, it is clear that this study  
232 did not provide any evidence for or against any effects caused by the transgene or the expressed  
233 protein. However, one thing almost everybody agreed on in the end, is that the Pusztai affair is a  
234 prime example for how *not* to handle potentially troubling findings, and how essential it is for  
235 scientists to stay in contact with, and explain their work to, the general public.

236 Now how does this relate to the CaMV 35S promoter? Several anti-GE groups and activists  
237 immediately picked up this story and, for some reason, highlighted the CaMV 35S promoter as  
238 the potential culprit of the observed health effects<sup>43,55</sup>. Greenpeace released a statement saying,  
239 *'For all we know they might have been caused by the virus used to transfer the alien DNA to the*  
240 *potatoes. This is the same virus used in Monsanto's Roundup Ready soy that is available in*  
241 *markets around the world'*, clearly ignorant to the fact that the CaMV 35S promoter is not a  
242 virus, but a short stretch of DNA, and the active ingredient in Roundup Ready Soy is also not a  
243 virus, but the EPSP synthase<sup>55</sup>. In this case, it was solely the origin of the CaMV 35S sequence  
244 from a virus that brought it to the attention of these groups. The word 'virus' certainly has a



245 negative connotation in most people’s mind, and thus seemed to be a good way to activate as  
246 many people against GE as possible. And other activists published work along similar lines<sup>43</sup>. So  
247 overall, the Pusztai study was simply a badly planned and poorly executed work, which most  
248 likely would have been significantly improved during a peer-review process, if some of the  
249 results had not been prematurely broadcast publicly on TV. But the poor handling by the people  
250 involved, in combination with a scientific community that failed to sufficiently inform and  
251 educate the public about genetic engineering and genetically modified organisms in general for  
252 over a decade, allowed this to escalate into an affair that shifted public perception of genetic  
253 engineering and genetically modified crops from healthy criticism to outright rejection.  
254 Unfortunately, the damage done could not be rectified to this day. Furthermore, due to the  
255 continuing lobbying of anti-GE activists, the reputation of the CaMV 35S promoter was also  
256 severely tarnished by this event, and it has since become one of the buzzwords of the GE-  
257 movement. This is probably best illustrated by a 2009 paper from famed anti-GE activist and  
258 pseudo-scientist Mae-Wan Ho, who claimed that eating transgenic crops carrying the CaMV 35S  
259 promoter could promote HIV in humans<sup>68</sup>. This claim is ‘backed up’ in the paper by a truly  
260 amazing line of argumentation: “*In humans, P-TEFb is required by HIV-1 for its transcription*  
261 *and replication. The long terminal repeat of HIV-1 has minimal promoter activity in the absence*  
262 *of the viral Tat protein. The CaMV 35S promoter, on the other hand, is strongly active in plant*  
263 *cells in the absence of any viral protein. Thus, the presence of CaMV 35S promoter effectively*  
264 *facilitates the transcription of HIV and other viruses*”<sup>68</sup>.

### 265 **The CaMV 35S Promoter Today (2000-today)**

266 To this day the CaMV 35S promoter remains the most commonly used promoter in plant science.  
267 Nonetheless, use of the CaMV 35S promoter is slowly decreasing due to several reasons. The  
268 number one reason being that today, in contrast to the 1980s, many alternatives to the CaMV 35S  
269 promoter are available to researchers. In academia, the *Arabidopsis UBIQUITIN10* promoter was  
270 identified in the mid-1990s as a strong promoter, active in all tissues of the plant body – indeed,  
271 the two major selling points of the CaMV 35S promoter – and was ready to replace it as a plant-  
272 derived promoter to use in plants<sup>69,70</sup>. By that time researchers had also discovered that the 35S  
273 promoter was actually not active in all tissues and cell types, but sometimes exhibited a ‘patchy’  
274 pattern, something not seen for the UBQ1 or 10 promoters<sup>71</sup>. Furthermore, by the year **2000**, the  
275 *Arabidopsis* genome had been sequenced and annotated, uncovering all genes and their putative

276 regulatory sequences<sup>72</sup>. Such sequences could now be easily cloned and enabled scientists to  
277 express their genes of interest under control of their respective native promoters, or very targeted  
278 in specific cells and tissues, and at physiological concentration levels. Gene silencing, which  
279 caused the problem in the Petunia field trial, prompted scientists to employ the 35S promoter in  
280 conjunction with the p19 silencing suppressor, further complicating the applicability of this  
281 promoter<sup>73</sup>. Finally, reports emerged that the 35S promoter could affect the expression not only  
282 of the downstream transgene, but also other genes in its vicinity, possibly via its enhancer  
283 regions, which further confounded the use of the 35S promoter<sup>74-76</sup>. Thus, while the CaMV 35S  
284 promoter is still used heavily in scientific studies, many now favor the use of endogenous  
285 promoters and/or the *UBIQUITIN10* promoter.

286 In agriculture, over 80 % of GE-crops in the field still carry a version of the CaMV 35S  
287 promoter, among those the most widely farmed varieties such as the Roundup Ready soybean, *Bt*  
288 corn and cotton, and the ‘Sunset’ papaya resistant to the papaya ringspot virus<sup>77-80</sup>. These crops  
289 have been found to be safe by all the major scientific institutions, and have been consumed by  
290 humans and livestock for decades now, without any negative health effects<sup>81-86</sup>. Since the  
291 generation, subsequent field-trials, safety tests and governmental approvals of a transgenic crop  
292 line are arduous, time-consuming and expensive, a switch to a different standard promoter will be  
293 a long-term project. The use of the CaMV 35S promoter is, however, limited because of multiple  
294 overlapping patents on it<sup>87</sup>. Among other things, this has led to enhanced use of similar, or related  
295 promoters, such as the figwort mosaic virus 34S promoter (FMV 34S)<sup>87,88</sup>. It was also found that  
296 in monocots, such as rice and corn, the CaMV 35S is not as active as it is in dicots, leading  
297 researchers to switch to, e.g., the later discovered *rice actin 1* or *maize Ubi-1* promoters<sup>89-92</sup>. So  
298 even though the switch is happening at a slower pace in applied agriculture than it is in academia,  
299 the variety of promoters used is steadily increasing also in this area.

300 Nonetheless, since its description in 1985 the CaMV 35S promoter has been the standard  
301 promoter used in all plant science and plant biotechnology, and has certainly propelled the  
302 research field forward like hardly any other discovery.

303

304

305 **Acknowledgements**

306 Thanks to Staffan Persson, Imre E. Somssich, Edward P. Rybicki, Raymond D. Shillito, and the  
307 Deutsche Forschungsgemeinschaft (German Research Foundation; Project 344523413) for  
308 comments and support.

309 **References**

- 310 1. **Schultz ES.** A transmissible mosaic disease of Chinese cabbage, mustard and turnip. **J**  
311 **Agric Res.** **1921**;22: 173–177.
- 312 2. **Larson RH, Walker JC.** A mosaic disease of cabbage. **J Agric Res.** **1939**;59: 367–392.
- 313 3. **Tompkins CM.** A transmissible mosaic disease of cauliflower. **J Agric Res.** **1937**;55: 33–  
314 46. Available: <https://naldc.nal.usda.gov/download/IND43968984/PDF>
- 315 4. **Broadbent L.** Investigation of Virus Diseases of Brassica Crops. **Agric Res Counc Rep**  
316 **Ser. Cambridge University Press; 1957**;: 94–ff. Available:  
317 <https://books.google.com.au/books?id=aIS2CgAAQBAJ>
- 318 5. **Day MF, Venables DG.** The Transmission of Cauliflower Mosaic Virus by Aphids. **Aust**  
319 **J Biol Sci.** **1961**;14: 187. Available at doi:10.1071/BI9610187
- 320 6. **Uzest M, Gargani D, Drucker M, Hébrard E, Garzo E, Candresse T, et al.** A protein  
321 key to plant virus transmission at the tip of the insect vector stylet. **Proc Natl Acad Sci U**  
322 **S A.** **2007**;104: 17959–64. Available at doi:10.1073/pnas.0706608104
- 323 7. **Shepherd RJ, Wakeman RJ, Romanko RR.** DNA in cauliflower mosaic virus.  
324 **Virology.** **1968**;36: 150–152. Available at doi:10.1016/0042-6822(68)90127-X
- 325 8. **Schultze M, Jiricny J, Hohn T.** Open Reading Frame-Viii Is Not Required for Viability  
326 of Cauliflower Mosaic-Virus. **Virology.** **1990**;176: 662–664.
- 327 9. **Franck A, Guilley H, Jonard G, Richards KE, Hirth L.** Nucleotide sequence of  
328 cauliflower mosaic virus DNA. **Cell.** **1980**;21: 285–94. Available at doi:10.1016/0092-  
329 8674(80)90136-1
- 330 10. **Hohn T, Hohn B, Lesot A, Lebeurier G.** Restriction map of native and cloned

- 331 cauliflower mosaic virus DNA. **Gene**. **1980**;11: 21–31.
- 332 11. **Covey SN, Hull R**. Transcription of cauliflower mosaic virus DNA. Detection of  
333 transcripts, properties, and location of the gene encoding the virus inclusion body protein.  
334 **Virology**. **1981**;111: 463–74. Available at doi:10.1016/0042-6822(81)90349-4
- 335 12. **Love AJ, Laird J, Holt J, Hamilton AJ, Sadanandom A, Milner JJ**. Cauliflower  
336 mosaic virus protein P6 is a suppressor of RNA silencing. **J Gen Virol**. **2007**;88: 3439–44.  
337 Available at doi:10.1099/vir.0.83090-0
- 338 13. **Kiss-László Z, Blanc S, Hohn T**. Splicing of cauliflower mosaic virus 35S RNA is  
339 essential for viral infectivity. **EMBO J**. **1995**;14: 3552–62. Available at  
340 doi:10.1002/j.1460-2075.1995.tb07361.x
- 341 14. **Pfeiffer P, Hohn T**. Involvement of reverse transcription in the replication of cauliflower  
342 mosaic virus: a detailed model and test of some aspects. **Cell**. **1983**;33: 781–9. Available  
343 at doi:10.1016/0092-8674(83)90020-X
- 344 15. **Guilley H, Dudley RK, Jonard G, Balázs E, Richards KE**. Transcription of Cauliflower  
345 mosaic virus DNA: detection of promoter sequences, and characterization of transcripts.  
346 **Cell**. **1982**;30: 763–73. Available:  
347 [http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Retrieve&db=PubMed&dopt=Citation](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Retrieve&db=PubMed&dopt=Citation&list_uids=7139714)  
348 [n&list\\_uids=7139714](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Retrieve&db=PubMed&dopt=Citation&list_uids=7139714)
- 349 16. **Blevins T, Rajeswaran R, Aregger M, Borah BK, Schepetilnikov M, Baerlocher L, et**  
350 **al**. Massive production of small RNAs from a non-coding region of Cauliflower mosaic  
351 virus in plant defense and viral counter-defense. **Nucleic Acids Res**. **2011**;39: 5003–14.  
352 Available at doi:10.1093/nar/gkr119
- 353 17. **Meyerowitz EM, Pruitt RE**. Arabidopsis thaliana and Plant Molecular Genetics. **Science**.  
354 **1985**;229: 1214–8. Available at doi:10.1126/science.229.4719.1214
- 355 18. **Somssich M**. A short history of Arabidopsis thaliana (L.) Heynh. Columbia-0. **PeerJ**  
356 **Prepr**. **2018**;e26931v3: 1–7. Available at doi:10.7287/peerj.preprints.26931v3
- 357 19. **Koncz C, De Greve H, André D, Deboeck F, Montagu MCE van, Schell J**. The opine

- 358 synthase genes carried by Ti plasmids contain all signals necessary for expression in  
359 plants. **EMBO J.** **1983**;2: 1597–603. Available at doi:10.1002/j.1460-  
360 2075.1983.tb01630.x
- 361 20. **Cashmore AR.** Nuclear Genes Encoding the Small Subunit of Ribulose-1,5-Bisphosphate  
362 Carboxylase. Genetic Engineering of Plants. Boston, MA: **Springer US**; **1983**. pp. 29–38.  
363 Available at doi:10.1007/978-1-4684-4544-2\_5
- 364 21. **Hull R.** The possible use of plant viral DNAs in genetic manipulation in plants. **Trends**  
365 **Biochem Sci.** **1978**;3: 254–256. Available at doi:10.1016/S0968-0004(78)95435-X
- 366 22. **Hohn B, Hohn T.** Cauliflower Mosaic Virus: A Potential Vector for Plant Genetic  
367 Engineering. Molecular Biology of Plant Tumors. **Elsevier**; **1982**. pp. 549–560. Available  
368 at doi:10.1016/B978-0-12-394380-4.50028-1
- 369 23. **Brisson N, Paszkowski J, Penswick JR, Gronenborn B, Potrykus I, Hohn T.**  
370 Expression of a bacterial gene in plants by using a viral vector. **Nature.** **1984**;310: 511–  
371 514. Available at doi:10.1038/310511a0
- 372 24. **Lefebvre DD, Miki BL, Laliberté J-F.** Mammalian Metallothionein Functions in Plants.  
373 Nature biotechnology. **1987**. pp. 32–34. Available at doi:10.1007/978-94-009-4482-4\_7
- 374 25. **Paszkowski J, Pisan B, Shillito RD, Hohn T, Hohn B, Potrykus I.** Genetic  
375 transformation of Brassica campestris var. rapa protoplasts with an engineered cauliflower  
376 mosaic virus genome. **Plant Mol Biol.** **1986**;6: 303–12. Available at  
377 doi:10.1007/BF00034937
- 378 26. **Somssich M.** A Short History of Plant Transformation. **PeerJ Prepr.** **2019**;: 1–28.  
379 Available at doi:10.7287/peerj.preprints.27556v1
- 380 27. **Haas M, Bureau M, Geldreich A, Yot P, Keller M.** Cauliflower mosaic virus: still in the  
381 news. **Mol Plant Pathol.** **2002**;3: 419–29. Available at doi:10.1046/j.1364-  
382 3703.2002.00136.x
- 383 28. **Gronenborn B, Gardner RC, Schaefer S, Shepherd RJ.** Propagation of foreign DNA in  
384 plants using cauliflower mosaic virus as vector. **Nature.** **1981**;294: 773–776. Available at

- 385           doi:10.1038/294773a0
- 386   29.   **Odell JT, Nagy F, Chua N-H.** Identification of DNA sequences required for activity of  
387       the cauliflower mosaic virus 35S promoter. **Nature.** **1985**;313: 810–2. Available at  
388       doi:10.1038/313810a0
- 389   30.   **Benfey PN, Chua N-H.** The Cauliflower Mosaic Virus 35S Promoter: Combinatorial  
390       Regulation of Transcription in Plants. **Science.** **1990**;250: 959–966. Available at  
391       doi:10.1126/science.250.4983.959
- 392   31.   **Fang R-X, Nagy F, Sivasubramaniam S, Chua N.** Multiple cis regulatory elements for  
393       maximal expression of the cauliflower mosaic virus 35S promoter in transgenic plants.  
394       **Plant Cell.** **1989**;1: 141–50. Available at doi:10.1105/tpc.1.1.141
- 395   32.   **Kay R, Chan A, Daly M, McPherson J.** Duplication of CaMV 35S Promoter Sequences  
396       Creates a Strong Enhancer for Plant Genes. **Science.** **1987**;236: 1299–302. Available at  
397       doi:10.1126/science.236.4806.1299
- 398   33.   **Ulmasov T, Murfett J, Hagen G, Guilfoyle TJ.** Aux/IAA Proteins Repress Expression of  
399       Reporter Genes Containing Natural and Highly Active Synthetic Auxin Response  
400       Elements. **Plant Cell.** **1997**;9: 1963–1971. Available at doi:10.1105/tpc.9.11.1963
- 401   34.   **Rushton PJ, Reinstädler A, Lipka V, Lippok B, Somssich IE.** Synthetic plant  
402       promoters containing defined regulatory elements provide novel insights into pathogen-  
403       and wound-induced signaling. **Plant Cell.** **2002**;14: 749–762. Available at  
404       doi:10.1105/tpc.010412
- 405   35.   **Zuo J, Niu Q-W, Chua N-H.** An estrogen receptor-based transactivator XVE mediates  
406       highly inducible gene expression in transgenic plants. **Plant J.** **2000**;24: 265–273.  
407       Available at doi:10.1046/j.1365-313x.2000.00868.x
- 408   36.   **Aoyama T, Chua N-H.** A glucocorticoid-mediated transcriptional induction system in  
409       transgenic plants. **Plant J.** **1997**;11: 605–12. Available at doi:10.1046/j.1365-  
410       313X.1997.11030605.x
- 411   37.   **Caddick MX, Greenland AJ, Jepson L, Krause K-P, Qu N, Riddell K V., et al.** An

- 412 ethanol inducible gene switch for plants used to manipulate carbon metabolism. **Nat**  
413 **Biotechnol.** **1998**;16: 177–180. Available at doi:10.1038/nbt0298-177
- 414 38. **Shah DM, Horsch RB, Klee HJ, Kishore GM, Winter JA, Tumer NE, et al.**  
415 Engineering Herbicide Tolerance in Transgenic Plants. **Science.** **1986**;233: 478–481.  
416 Available at doi:10.1126/science.233.4762.478
- 417 39. **Koshiba T.** Shikimate Kinase and 5-Enolpyruvylshikimate-3-phosphate Synthase in  
418 Phaseolus mungo Seedlings. **Zeitschrift für Pflanzenphysiologie. Gustav Fischer**  
419 **Verlag, Stuttgart; 1978**;88: 353–355. Available at doi:10.1016/S0044-328X(78)80138-X
- 420 40. **Steinrücken HC, Amrhein N.** The herbicide glyphosate is a potent inhibitor of 5-  
421 enolpyruvyl-shikimic acid-3-phosphate synthase. **Biochem Biophys Res Commun.**  
422 **1980**;94: 1207–12. Available at doi:10.1016/0006-291X(80)90547-1
- 423 41. **Herrera-Estrella L, Depicker A, Montagu MCE van, Schell J.** Expression of chimaeric  
424 genes transferred into plant cells using a Ti-plasmid-derived vector. **Nature.** **1983**;303:  
425 209–213. Available at doi:10.1038/303209a0
- 426 42. **Hull R, Covey SN, Dale P.** Genetically modified plants and the 35S promoter: assessing  
427 the risks and enhancing the debate. **Microb Ecol Health Dis.** **2000**;12: 1–5. Available at  
428 doi:10.1080/089106000435527
- 429 43. **Ho M-W, Ryan A, Cummins J.** Cauliflower Mosaic Viral Promoter - A Recipe for  
430 Disaster? **Microb Ecol Health Dis.** **1999**;11: 194–197. Available at  
431 doi:10.1080/08910609908540827
- 432 44. **Lange M.** Erster Freilandversuch mit gentechnisch manipulierten Pflanzen.  
433 **Deutschlandfunk.** **2015**;: 18–20. Available: [http://www.deutschlandfunk.de/koeln-erster-](http://www.deutschlandfunk.de/koeln-erster-freilandversuch-mit-gentechnisch-manipulierten.871.de.html?dram:article_id=319729)  
434 [freilandversuch-mit-gentechnisch-manipulierten.871.de.html?dram:article\\_id=319729](http://www.deutschlandfunk.de/koeln-erster-freilandversuch-mit-gentechnisch-manipulierten.871.de.html?dram:article_id=319729)
- 435 45. **Randerson J.** Arpad Pusztai: Biological divide. **Guard.** **2008**;: 18–23. Available:  
436 <http://www.guardian.co.uk/education/2008/jan/15/academicexperts.highereducationprofile>
- 437 46. **Meyer P, Heidmann I, Forkmann G, Saedler H.** A new petunia flower colour generated  
438 by transformation of a mutant with a maize gene. **Nature.** **1987**;330: 677–8. Available at

- 439 doi:10.1038/330677a0
- 440 47. **McClintock B.** The origin and behavior of mutable loci in maize. **Proc Natl Acad Sci U S**  
441 **A.** **1950**;36: 344–55. Available at doi:10.1073/pnas.36.6.344
- 442 48. **Meyer P, Linn F, Heidmann I, Meyer z. A. H, Niedenhof I, Saedler H.** Endogenous  
443 and environmental factors influence 35S promoter methylation of a maize A1 gene  
444 construct in transgenic petunia and its colour phenotype. **Mol Gen Genet.** **1992**;231: 345–  
445 52. Available at doi:10.1007/BF00292701
- 446 49. **Linn F, Heidmann I, Saedler H, Meyer P.** Epigenetic changes in the expression of the  
447 maize A1 gene in *Petunia hybrida*: role of numbers of integrated gene copies and state of  
448 methylation. **Mol Gen Genet.** **1990**;222: 329–36. Available at doi:10.1007/BF00633837
- 449 50. **Specter M.** West Germany's anguished science. **Washington Post.** **1990**;April 11: 1–4.  
450 Available: [https://www.washingtonpost.com/archive/politics/1990/04/11/west-germanys-](https://www.washingtonpost.com/archive/politics/1990/04/11/west-germanys-anguished-science/41026c6c-c574-46bc-88c2-45491f9daa2d/?utm_term=.1f45ce3efb2a)  
451 [anguished-science/41026c6c-c574-46bc-88c2-45491f9daa2d/?utm\\_term=.1f45ce3efb2a](https://www.washingtonpost.com/archive/politics/1990/04/11/west-germanys-anguished-science/41026c6c-c574-46bc-88c2-45491f9daa2d/?utm_term=.1f45ce3efb2a)
- 452 51. **Billstein H.** Ein Zaun schützt die Petunien. **Zeit.** **1988**;46: 7–10. Available:  
453 <http://www.zeit.de/1988/46/ein-zaun-schuetzt-die-petunien>
- 454 52. **Federal Ministry for the Environment; Nature Conservation; Building and Nuclear**  
455 **Safety.** 2015 Nature Awareness Study - Population survey on nature and biological  
456 diversity. 2015 Nature Awareness Study. Bonn; **2015.** Available:  
457 <https://www.bfn.de/themen/gesellschaft/naturbewusstsein/studie-2015.html>
- 458 53. **Servick K.** How the transgenic petunia carnage of 2017 began. **Science.** **2017**;: 0–3.  
459 Available at doi:10.1126/science.aan6886
- 460 54. **Haselmair-Gosch C, Miosic S, Nitarska D, Roth BL, Walliser B, Paltram R, et al.**  
461 Great Cause—Small Effect: Undeclared Genetically Engineered Orange Petunias Harbor  
462 an Inefficient Dihydroflavonol 4-Reductase. **Front Plant Sci.** **2018**;9: 1–12. Available at  
463 doi:10.3389/fpls.2018.00149
- 464 55. **Institute for Agriculture & Trade Policy.** Greenpeace Calls for Immediate Total Ban on  
465 GMO Food. **Inst Agric Trade Policy.** **1999**;: 18–20. Available:



- 466 <https://www.iatp.org/news/greenpeace-calls-for-immediate-total-ban-on-gmo-food>
- 467 56. **Enserink M.** Institute Copes With Genetic Hot Potato. **Science**. **1998**;281: 1124b–1125.  
468 Available at doi:10.1126/science.281.5380.1124b
- 469 57. **Fedoroff N V., Brown N.** Mendel in the Kitchen, a Scientist’s View of Genetically  
470 Modified Food. Mendel in the Kitchen, a Scientist’s View of Genetically Modified Food.  
471 **Joseph Henry Press; 2004.**
- 472 58. **Enserink M.** Preliminary Data Touch Off Genetic Food Fight. **Science**. **1999**;283: 1094–  
473 1095. Available at doi:10.1126/science.283.5405.1094
- 474 59. **Ewen SWB, Pusztai Á.** Effect of diets containing genetically modified potatoes  
475 expressing Galanthus nivalis lectin on rat small intestine. **Lancet**. **1999**;354: 1353–4.  
476 Available at doi:10.1016/S0140-6736(98)05860-7
- 477 60. **Macedo MLR, Oliveira CFR, Oliveira CT.** Insecticidal activity of plant lectins and  
478 potential application in crop protection. **Molecules**. **2015**;20: 2014–33. Available at  
479 doi:10.3390/molecules20022014
- 480 61. **Du J, Foissac X, Carss A, Gatehouse AMR, Gatehouse JA.** Ferritin acts as the most  
481 abundant binding protein for snowdrop lectin in the midgut of rice brown planthoppers  
482 (*Nilaparvata lugens*). **Insect Biochem Mol Biol**. **2000**;30: 297–305. Available at  
483 doi:10.1016/S0965-1748(99)00130-7
- 484 62. **Pusztai Á, Ewen SWB, Grant G, Peumans WJ, Van Damme EJM, Rubio L, et al.**  
485 Relationship between Survival and Binding of Plant Lectins during Small Intestinal  
486 Passage and Their Effectiveness as Growth Factors. **Digestion**. **1990**;46: 308–316.  
487 Available at doi:10.1159/000200402
- 488 63. **Gatehouse AMR, Down RE, Powell KS, Sauvion N, Rahbé Y, Newell CA, et al.**  
489 Transgenic potato plants with enhanced resistance to the peach-potato aphid *Myzus*  
490 *persicae*. **Entomol Exp Appl**. **1996**;79: 295–307. Available at doi:10.1111/j.1570-  
491 7458.1996.tb00837.x
- 492 64. **Kuiper HA, Noteborn HP, Peijnenburg AA.** Adequacy of methods for testing the safety

- 493 of genetically modified foods. **Lancet**. **1999**;354: 1315–6. Available at  
494 doi:10.1016/S0140-6736(99)00341-4
- 495 65. **The Royal British Society**. Review of data on possible toxicity of GM potatoes. **R Br**  
496 **Soc**. **1999**;10: 391–399.
- 497 66. **Enserink M**. The Lancet scolded over Pusztai paper. **Science**. **1999**;286: 656. Available at  
498 doi:10.1126/science.286.5440.656a
- 499 67. **Down RE, Ford L, Bedford SJ, Gatehouse LN, Newell C, Gatehouse JA, et al**.  
500 Influence of plant development and environment on transgene expression in potato and  
501 consequences for insect resistance. **Transgenic Res**. **2001**;10: 223–36. Available:  
502 [http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Retrieve&db=PubMed&dopt=Citation&list\\_uids=11437279](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Retrieve&db=PubMed&dopt=Citation&list_uids=11437279)  
503
- 504 68. **Ho M-W, Cummins J**. New evidence links CaMV 35S promoter to HIV transcription.  
505 **Microb Ecol Health Dis**. **2009**;21: 172–174. Available at  
506 doi:10.3109/08910600903495053
- 507 69. **Sun CW, Callis J**. Independent modulation of Arabidopsis thaliana polyubiquitin mRNAs  
508 in different organs and in response to environmental changes. **Plant J**. **1997**;11: 1017–27.  
509 Available at doi:10.1046/j.1365-313X.1997.11051017.x
- 510 70. **Norris SR, Meyer SE, Callis J**. The intron of Arabidopsis thaliana polyubiquitin genes is  
511 conserved in location and is a quantitative determinant of chimeric gene expression. **Plant**  
512 **Mol Biol**. **1993**;21: 895–906. Available at doi:10.1007/BF00027120
- 513 71. **Holtorf S, Apel K, Bohlmann H**. Comparison of different constitutive and inducible  
514 promoters for the overexpression of transgenes in Arabidopsis thaliana. **Plant Mol Biol**.  
515 **1995**;29: 637–46. Available at doi:10.1007/BF00041155
- 516 72. **Arabidopsis Genome Initiative**. Analysis of the genome sequence of the flowering plant  
517 Arabidopsis thaliana. **Nature**. **2000**;408: 796–815. Available at doi:10.1038/35048692
- 518 73. **Van Der Hoorn RAL, Rivas S, Wulff BBH, Jones JDG, Joosten MHAJ**. Rapid  
519 migration in gel filtration of the Cf-4 and Cf-9 resistance proteins is an intrinsic property

- 520 of Cf proteins and not because of their association with high-molecular-weight proteins.  
521 **Plant J.** **2003**;35: 305–15. Available at doi:10.1046/j.1365-313X.2003.01803.x
- 522 74. **Yoo SY, Bomblies K, Yoo SK, Yang JW, Choi MS, Lee JS, et al.** The 35S promoter  
523 used in a selectable marker gene of a plant transformation vector affects the expression of  
524 the transgene. **Planta.** **2005**;221: 523–30. Available at doi:10.1007/s00425-004-1466-4
- 525 75. **Weigel D, Ahn JH, Blázquez MA, Borevitz JO, Christensen SK, Fankhauser C, et al.**  
526 Activation Tagging in Arabidopsis. **Plant Physiol.** **2000**;122: 1003–1013. Available at  
527 doi:10.1104/pp.122.4.1003
- 528 76. **Gudynaite-Savitch L, Johnson DA, Miki BLA.** Strategies to mitigate transgene-  
529 promoter interactions. **Plant Biotechnol J.** **2009**;7: 472–85. Available at  
530 doi:10.1111/j.1467-7652.2009.00416.x
- 531 77. **U.S. Department of Agriculture.** USDA/APHIS Petition 96-051-01P for the  
532 Determination of Nonregulated Status for Transgenic “Sunset” Papaya Lines 55-1 and 63-  
533 1. **Fed Regist.** **1996**;61: 1–23. Available:  
534 [https://www.aphis.usda.gov/brs/aphisdocs2/96\\_05101p\\_com.pdf](https://www.aphis.usda.gov/brs/aphisdocs2/96_05101p_com.pdf)
- 535 78. **U.S. Environmental Protection Agency.** Cry1Ab and Cry1F Bacillus thuringiensis (Bt)  
536 Corn Plant-Incorporated Protectants. **Biopestic Regist ACTION Doc.** **2010**; Available:  
537 [https://web.archive.org/web/20130127095310/http://www.epa.gov/oppbppd1/biopesticides](https://web.archive.org/web/20130127095310/http://www.epa.gov/oppbppd1/biopesticides/pips/cry1f-cry1ab-brad.pdf)  
538 [/pips/cry1f-cry1ab-brad.pdf](https://web.archive.org/web/20130127095310/http://www.epa.gov/oppbppd1/biopesticides/pips/cry1f-cry1ab-brad.pdf)
- 539 79. **Perlak FJ, Deaton RW, Armstrong TA, Fuchs RL, Sims SR, Greenplate JT, et al.**  
540 Insect resistant cotton plants. **Bio/technology.** **1990**;8: 939–43. Available at  
541 doi:10.1038/nbt1090-939
- 542 80. **Padgett SR, Kolacz KH, Delannay X, Re DB, LaVallee BJ, Tinius CN, et al.**  
543 Development, Identification, and Characterization of a Glyphosate-Tolerant Soybean Line.  
544 **Crop Sci.** **1995**;35: 1451. Available at  
545 doi:10.2135/cropsci1995.0011183X003500050032x
- 546 81. **Norero D.** More than 280 scientific and technical institutions support the safety of GM  
547 crops. **Chile Bio.** **2014**; Available: <http://www.siquierotransgenicos.cl/2015/06/13/more->

- 548 than-240-organizations-and-scientific-institutions-support-the-safety-of-gm-crops/
- 549 82. **AAAS Board of Directors**. Statement by the AAAS Board of Directors On Labeling of  
550 Genetically Modified Foods. **Am Assoc Adv Sci**. **2012**;: 1. Available:  
551 <https://www.aaas.org/news/statement-aaas-board-directors-labeling-genetically-modified->  
552 [foods](https://www.aaas.org/news/statement-aaas-board-directors-labeling-genetically-modified-foods)
- 553 83. **European Commission**. A decade of EU-funded GMO research (2001–2010).  
554 **Research\*eu**. **2010**;: 1–264. Available at doi:doi 10.2777/97784
- 555 84. **Pellegrino E, Bedini S, Nuti M, Ercoli L**. Impact of genetically engineered maize on  
556 agronomic, environmental and toxicological traits: a meta-analysis of 21 years of field  
557 data. **Sci Rep. Springer US**; **2018**;8: 3113. Available at doi:10.1038/s41598-018-21284-2
- 558 85. **Yi D, Fang Z, Yang L**. Effects of Bt cabbage pollen on the honeybee *Apis mellifera* L.  
559 **Sci Rep. Springer US**; **2018**;8: 482. Available at doi:10.1038/s41598-017-18883-w
- 560 86. **Snell C, Bernheim A, Bergé J-B, Kuntz M, Pascal G, Paris A, et al**. Assessment of the  
561 health impact of GM plant diets in long-term and multigenerational animal feeding trials:  
562 A literature review. **Food Chem Toxicol. Elsevier Ltd**; **2012**;50: 1134–1148. Available at  
563 doi:10.1016/j.fct.2011.11.048
- 564 87. **Chi-Ham CL, Boettiger S, Figueroa-Balderas R, Bird S, Geoola JN, Zamora P, et al**.  
565 An intellectual property sharing initiative in agricultural biotechnology: development of  
566 broadly accessible technologies for plant transformation. **Plant Biotechnol J**. **2012**;10:  
567 501–10. Available at doi:10.1111/j.1467-7652.2011.00674.x
- 568 88. **Graff GD, Cullen SE, Bradford KJ, Zilberman D, Bennett AB**. The public-private  
569 structure of intellectual property ownership in agricultural biotechnology. **Nat Biotechnol**.  
570 **2003**;21: 989–95. Available at doi:10.1038/nbt0903-989
- 571 89. **Park S-H, Yi N, Kim YS, Jeong M-H, Bang S-W, Choi Y Do, et al**. Analysis of five  
572 novel putative constitutive gene promoters in transgenic rice plants. **J Exp Bot**. **2010**;61:  
573 2459–67. Available at doi:10.1093/jxb/erq076
- 574 90. **Christensen AH, Sharrock RA, Quail PH**. Maize polyubiquitin genes: structure, thermal

575 perturbation of expression and transcript splicing, and promoter activity following transfer  
576 to protoplasts by electroporation. **Plant Mol Biol.** **1992**;18: 675–89. Available at  
577 doi:10.1007/BF00020010

578 91. **McElroy D, Zhang W, Cao J, Wu R.** Isolation of an Efficient Actin Promoter for Use in  
579 Rice Transformation. **Plant Cell.** **1990**;2: 163–171. Available at doi:10.1105/tpc.2.2.163

580 92. **Nuccio ML.** A Brief History of Promoter Development for Use in Transgenic Maize  
581 Applications. *Methods in Molecular Biology.* **2018.** pp. 61–93. Available at  
582 doi:10.1007/978-1-4939-7315-6\_4

583