**Supplementary File 4. Plasmids employed in this study**

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| Plasmid | Description | Reference |
| pBluescript SK(+) | Cloning vector, *amp*R | Thermo Scientific |
| pME4305 | *six:*P*xylP:β-rec:trpC*t*:phleo*R*:six* | (Thieme et al., 2018) |
| pME4319 | *Swa*I*:six:*P*xylP:β-rec:trpC*t*:phleo*R*:six:Pml*I | This study |
| pME4574 | *5’UTR(veA): nat*RM*:3’UTR(veA)* | (Thieme et al., 2018) |
| pME4605 | *5’UTR(velB): nat*RM*:3’UTR(velB)* | (Thieme et al., 2018) |
| pME4636 | *5’UTR(laeA):phleoRM:3’UTR(laeA)* | This study |
| pME4645 | 5’UTR(*xptC*)*:xptC:gfp:phleo*RM*:*3’UTR(*xptC*) | This study |
| pME4842 | 5’UTR(*mdpG*)*:phleo*RM*:*3’UTR(*mdpG*) | This study |
| pME4843 | 5’UTR(*mdpF*)*:phleo*RM*:*3’UTR(*mdpF*) | This study |
| pME4844 | 5’UTR(*mdpC*)*:phleo*RM*:*3’UTR(*mdpC*) | This study |
| pME4845 | 5’UTR(*mdpL*)*:phleo*RM*:*3’UTR(*mdpL*) | This study |
| pME4846 | 5’UTR(*mdpD*)*:phleo*RM*:*3’UTR(*mdpD*) | This study |
| pME4847 | 5’UTR(*xptA*)*:phleo*RM*:*3’UTR(*xptA*) | This study |
| pME4848 | 5’UTR(*xptB*)*:phleo*RM*:*3’UTR(*xptB*) | This study |
| pME4849 | 5’UTR(*xptC*)*:phleo*RM*:*3’UTR(*xptC*) | This study |
| pME4850 | 5’UTR(*mdpG*)*:mdpG:phleo*RM*:*3’UTR(*mdpG*) | This study |
| pME4851 | 5’UTR(*mdpC*)*:mdpC:phleo*RM*:*3’UTR(*mdpC*) | This study |

P = promoter, t = terminator, R = resistance, *phleo*RM = recyclable phleomycin resistance cassette, *six* = β-recombinase recognition sequence

**References**

Thieme, K. G., Gerke, J., Sasse, C., Valerius, O., Thieme, S., Karimi, R., Heinrich, A. K., Finkernagel, F., Smith, K., & Bode, H. B. (2018). Velvet domain protein VosA represses the zinc cluster transcription factor SclB regulatory network for *Aspergillus nidulans* asexual development, oxidative stress response and secondary metabolism. *PLoS Genetics, 14*(7), e1007511. doi:<https://doi.org/10.1371/journal.pgen.1007511>