Trichoderma genes involved in sensing of the mating partner

Sensing of a potential mating partner is a prerequisite for sexual reproduction, and is fulfilled by a pheromone system consisting of peptide pheromone precursors and pheromone receptors (Nieuwenhuis & Aanen, 2012). Genome analysis shows that all 12 *Trichoderma* spp. have genes encoding the α -type peptide pheromone precursor PPG1, which is comparable to its homolog in other fungi.

	T. reesei	T. longibrachiatum	T. citrinoviride	T. parareesei	T. harzianum	T. guizhouense	T. afroharzianum	T. virens	T. atroviride	T. gamsii	T. asperellum	T. hamatum
PPG1, α type	104292	317925	1198762	OTA03422	86570	OPB47026	1241	44505	297240	18666200	144740	4957
HPP1	34493	1420161	1128304	*	463451	OPB39925	830	91991		*	*	•
STE2 = HPR2	64018	29766	1115433	OTA07042	96494	OPB41683	4616	21417	17079, 36032	18661910	44057	7747
STE3 = HPR1	57526	1322746	1207301	OTA04631	107686	OPB39349	6116	39157	147894	18660796	127205	1425
* no annotation, gene presence confirmed by translation of nt sequences												

In contrast, *Trichoderma* does not contain orthologs of the **a**-type peptide pheromone precursors, but instead has h-type precursors (HPP1), which share characteristics of **a**- and α -types by containing the consensus motif (LI)GC(TS)VM (Schmoll *et al.*, 2010). Interestingly, the HPP1 orthologs of members of ST are shorter because they contain only 1 or 2 copies of the consensus motif: In addition, all *Trichoderma* spp. contain the respective peptide pheromone precursors bind to pheromone receptors HPR1 (*ste3*) and HPR2 (*ste2*), which typical for heterothallic ascomycetes, all *Trichoderma* spp. have one MAT1-1 associated STE2p-type peptide pheromone receptor (HPR2), and one MAT1-2 associated STE3p-type peptide pheromone receptor (HPTR1).

T. atroviride M	SEGVQSFSAVQAKKGQSPQNSPATSQSVKTHFIGYLGCTIM GPGLLPTTTSGCTVQAKP-KPQQGSVKTHYVGFMGCTVM RSAQTVMTKPQGKPQYIGYLGCTVMAKPQDKPQGKPQYIGYLGCTVM RPSPASRPAQPTKGGYVGYLGCTVMAKPQKDGLGWTNSDCTVMAKPAKGGYVGYLGCTVM
T. gamsii 🔤	GPGLLPTTTSGCTVQAKE-KPQQGSVKTHYVGFMGCTVM
T. asperellum M	RSAQTVMTKPQGKPQYIGYL <mark>G</mark> CTVMAK <mark>P</mark> QDKPQGKPQ <mark>YIGYLGCTVM</mark>
T. hamatum 🍱	RPSPASRPAQPTKGGYVGYL <mark>C</mark> CTVMAK <mark>P</mark> QKDGLGWTNSDCTVMAKPAKGG <mark>YVGYLGCTVM</mark>

Sequence of the HPP1 proteins from Trichoderma spp. of ST. The conserved YVGYLGCTVM motif is underlined.

References

Nieuwenhuis BPS & Aanen DK (2012) Sexual selection in fungi. *J Evolution Biol* **25**: 2397-2411. Schmoll M, Seibel C, Tisch D, Dorrer M & Kubicek CP (2010) A novel class of peptide pheromone precursors in ascomycetous fungi. *Mol Microbiol* **77**: 1483-1501.