

Additional file 1: Table S1. Summary of RNA-sequencing data generated from *Penicillium oxalicum* strain HP7-1.

Samples^a	Clean Reads	Number of nucleotides (bp)	Overall read alignment rate (%)	Concordant pair alignment rate (%)	Number of expressed genes
HP7-1_WR_S-1	24298252	2429825200	87.62	72.80	8950
HP7-1_WR_S-2	23826040	2382604000	94.89	76.38	8721
HP7-1_WR_S-3	23783778	2378377800	94.88	76.67	8711
HP7-1_WR_L-1	23992762	2399276200	85.95	71.25	8850
HP7-1_WR_L-2	23811528	2381152800	89.70	73.15	8932
HP7-1_WR_L-3	23751856	2375185600	90.11	71.76	8830

^a WR: wheat bran plus rice straw; S: solid-state fermentation; L: submerged fermentation.