

Sample	Total Paired Reads	Mapped Reads
WT (1)	26,820,778	25,069,976 (93%)
WT (2)	26,799,612	24,896,074 (93%)
WT (3)	24,623,533	22,981,899 (93%)
<i>ppn1</i> Δ (1)	27,074,925	25,654,708 (95%)
<i>ppn1</i> Δ (2)	21,792,674	20,210,444 (93%)
<i>ppn1</i> Δ (3)	28,302,106	26,637,590 (94%)
<i>swd22</i> Δ (1)	24,435,616	23,082,133 (94%)
<i>swd22</i> Δ (2)	24,699,951	23,577,405 (95%)
<i>swd22</i> Δ (3)	23,445,504	22,211,684 (95%)
<i>dis2</i> Δ (1)	26,747,724	25,468,701 (95%)
<i>dis2</i> Δ (2)	23,488,584	22,144,361 (94%)
<i>dis2</i> Δ (3)	24,673,733	23,340,878 (95%)

S1 Fig. RNA-seq read counts for triplicate biological replicates.