

**Table S3 Introgression analysis.** Tests for introgression of the P3 strain into P1 or P2 strains that are more closely related with each other than to P1, using stickleback as the outgroup (O). D% is the introgression test statistic, on which the standard error (SE), 95% confidence intervals (CI) and Z-scores were estimated using block-wise jackknife ([Durand et al. 2011](#)).

P1	P2	P3	O	D%	SE	95% CI	Z-score
<b>Kaga</b>	<b>Hni</b>	<b>HdrR</b>	Stickleback	<b>3.6</b>	<b>1.3</b>	<b>2.5</b>	<b>2.77</b>
Kaga	Hni	Wild Southern	"	2.6	1	2	2.5
Kaga	Hni	Wild Southern (SNPs with HdrR)	"	-4	3.6	7.1	1.1
<b>Nilan</b>	<b>Hsok</b>	<b>HdrR</b>	"	<b>-12.3</b>	<b>4.66</b>	<b>9.1</b>	<b>2.64</b>
Nilan	Hsok	Wild Southern	"	-10.4	5.27	10.3	1.96
Nilan	Hsok	Wild Southern (SNPs with HdrR)	"	7.6	6.5	12.8	0.6
Nilan	Hsok	Kaga	"	-0.7	2.5	4.9	0.28

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