

Table S1. Overview of the whole-genome sequencing data

ID	Tissue type	Protocol	# Raw reads	# Mappable reads	# Clean reads*	Depth
ACC1-II-1	Blood	100bp, PE	3,310,551,412	3,153,459,526	2,707,907,255	90X
DS1-II-2	Blood	100bp, PE	3,225,109,320	3,097,723,302	2,393,004,900	80X
DS2-I-1	Blood	100bp, PE	2,990,509,808	2,871,412,207	2,276,650,271	76X

*: The number of reads after removing improperly-paired, duplicated, multiple-hit and too-many-error reads