

Additional file 2. Summary of mRNA-Seq read counts and mapping statistics

Individuals	NO. Reads	Genomic Reads		Junction Reads
		unique	non-unique	unique
PT02	58.9M	39.5M	4.9M	5.5M
PT03	51.2M	35.1M	4.7M	4.3M
PT04	66.5M	45.4M	7.3M	6.3M
PT05	61.3M	41.5M	8.1M	6.0M
PT06	59.9M	40.8M	5.8M	5.1M
PT07	60.0M	38.5M	5.2M	4.3M
PT08	52.1M	36.4M	4.2M	5.0M
PT09	62.8M	41.4M	6.1M	5.2M
PT10	90.4M	58.6M	9.4M	8.5M
PT11	66.7M	28.2M	6.5M	4.0M
PT12	106.4M	66.5M	10.5M	9.5M
PT13	121.8M	78.8M	13.4M	12.0M
PT14	96.9M	60.3M	9.9M	7.8M
PT15	110.5M	70.2M	11.1M	9.8M
PT16	98.0M	65.0M	10.6M	9.0M
PT17	100.1M	65.8M	9.7M	9.3M
PT18	94.8M	61.0M	9.3M	8.2M
PT19	100.3M	65.0M	10.1M	9.2M
PT20	106.4M	62.9M	10.5M	8.6M
PT21	102.5M	60.0M	10.2M	8.1M
Total	1668M	1061M	169M	146M
Fractions	100%	63.6%	10.1%	8.7%