Table S1. Primer-specific amplification efficiencies of the candidate reference gene.

Gene	Efficiency (%)	R ²		
ACTB	92.0	0.9966		
GAPDH	103.0	0.9948		
UXT	108.0	0.9987		
TBP	95.0	0.9972		
YWHAZ	93.0	0.9999		
RPL13A	92.0	0.9999		
SDHA	105.0	0.9999		
RPS15	103.0	1.0000		
HPRT1	106.0	1.0000		
PPIA	95.0	0.9900		
HMBS	100.0	0.9999		
MRPL39	93.0	0.9946		
PPP1R11	102.0	0.9996		
TET2	104.0	0.9994		

Table S2. Averaged raw Ct values obtained for each candidate gene in all testis samples.

	Sample Name 1	PPIA	RPS15	MRPL39	PPP1R11	SDHA	TBP	UXT	ACTB	HMBS	GAPDH	YWHAZ	HPRT1	RPL13A
	6M4-1	20.29	19.28	26.27	27.26	28.59	27.45	24.63	19.42	31.08	28.05	23.33	27.68	21.92
	6M6-1	22.31	19.18	26.41	26.41	27.66	27.42	25.63	20.61	31.23	27.64	22.66	28.23	22.00
	6M7-2	22.41	19.08	26.47	26.85	27.27	27.49	25.48	20.56	30.81	27.65	22.69	28.12	21.92
	18M2-1	21.16	19.21	25.08	26.82	27.00	26.87	25.24	20.60	30.22	26.69	22.30	27.76	22.38
	18M4-1	22.43	19.82	26.27	26.38	27.87	27.30	25.88	20.73	30.80	27.13	22.47	28.25	22.69
	18M5-1	21.01	20.19	24.60	25.60	26.94	26.08	25.33	20.41	29.75	26.91	21.69	26.36	22.79
	30M10-1	20.05	20.20	23.22	23.92	26.77	25.67	25.61	21.23	28.89	27.59	21.10	26.12	23.09
	30M7-1	20.64	20.65	23.66	23.96	27.27	25.75	25.74	20.82	28.86	27.81	21.43	26.47	23.49
	30M8-1	20.29	20.33	23.69	24.39	28.08	26.19	25.46	21.34	28.69	28.22	21.76	26.41	23.17
	6Y-3	19.84	20.05	23.07	23.29	27.28	25.44	25.71	20.88	28.50	28.07	20.62	26.39	22.87
	6Y-4	20.08	20.35	23.20	23.12	27.17	25.47	25.93	20.97	28.28	27.86	20.57	26.38	23.03
	6Y-6	20.74	21.23	24.07	24.32	28.67	26.69	26.57	22.20	30.07	29.41	22.04	26.92	24.22
All development	average	20.94	19.96	24.67	25.19	27.55	26.49	25.60	20.81	29.76	27.75	21.89	27.09	22.80
stages	SD	0.91	0.64	1.31	1.45	0.60	0.78	0.44	0.62	1.04	0.67	0.82	0.81	0.66
T	average	21.60	19.46	25.85	26.55	27.55	27.10	25.37	20.39	30.65	27.35	22.52	27.73	22.28
Immature stages	SD	0.83	0.40	0.73	0.52	0.57	0.50	0.39	0.44	0.51	0.47	0.49	0.65	0.36
3.6	average	20.27	20.47	23.49	23.83	27.54	25.87	25.84	21.24	28.88	28.16	21.25	26.45	23.31
Mature stages	SD	0.32	0.39	0.35	0.48	0.64	0.44	0.36	0.47	0.57	0.59	0.55	0.24	0.45

¹ Sample Name: 6M-6 months, 18M-18 months, 30M-30months, 6Y-6 years.

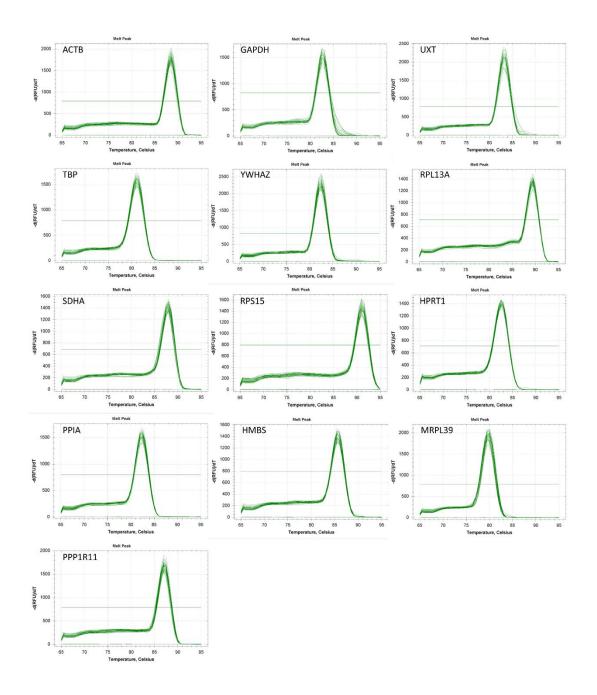


Figure S1. Melting curves of 13 genes of RT-qPCR.