

Table S1. *MTNR1A* gene amplified regions and used primers

Amplified regions	Forward primers	Reverse primers
Promoter	GCACAAAAAGAACCCAAGGA	TCAGGTGTCGCACTGTAACC
Promoter	TGTTCCCTGGAGAAAGTCTGG	CATGCATCAACACTGGACTG
Exon 1	CTCGACGCTCTGGGGAT	CCGAACAGGGAAGAGGTTG
Exon 2 and 3'UTR	GGCCCTAACCATGTTTCT	CTCCCACTCTGTTCCCTGAA

Table S2. PCR conditions for *MTNR1A* studied regions

PCR steps	Promoter		Promoter		Exon 1		Exon 2 and 3' UTR	
	Initial denaturation	94°C 3min	94°C 3min					
35 cycles	Denaturation	94°C 1min	94°C 1min					
	Annealing	60°C 1min	60°C 1min	50°C 1min	54°C 1min	54°C 1min	54°C 1min	54°C 1min
	Extension	72°C 1min	72°C 1min	72°C 30sec	72°C 1min	72°C 1min	72°C 1min	72°C 1min
Final elongation		72°C 10min	72°C 10min					

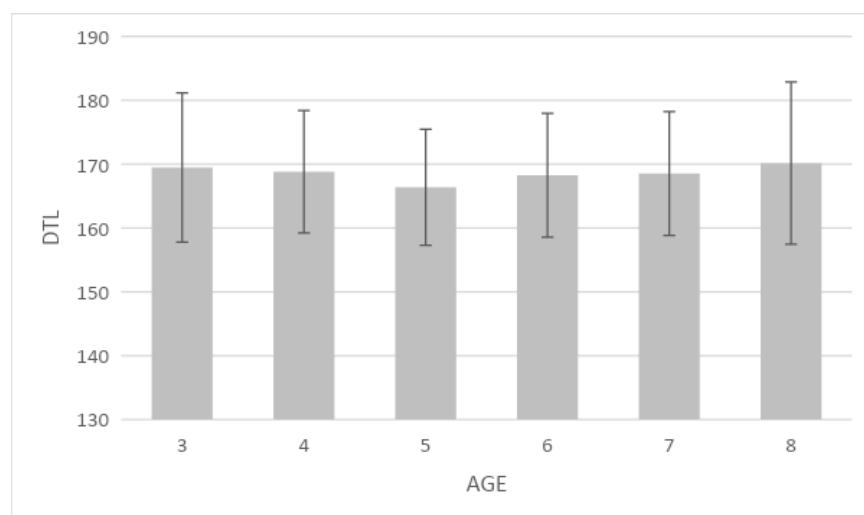


Figure S1. Graph showing the mean and the standard deviation of DTL divided by age, in the studied Lebanese Awassi population (n=165). DTL = Days To Lambing

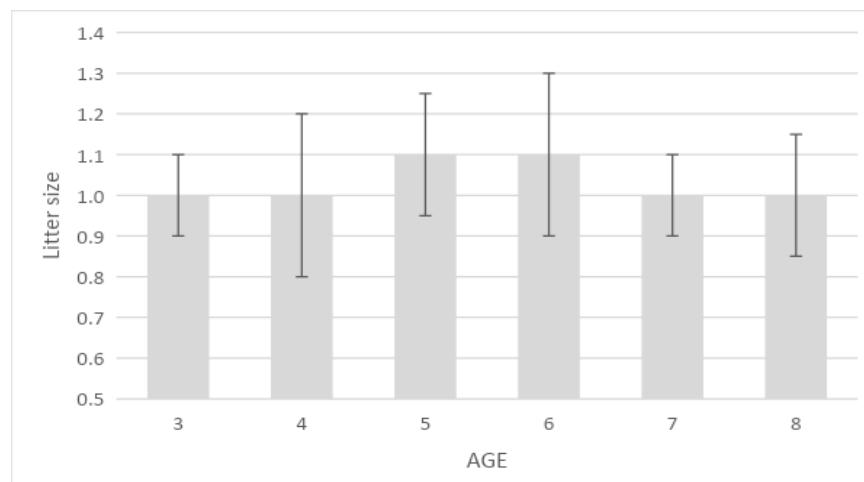


Figure S2. Graph showing the mean and the standard deviation of litter size divided by age, in the studied Lebanese Awassi population (n=165).

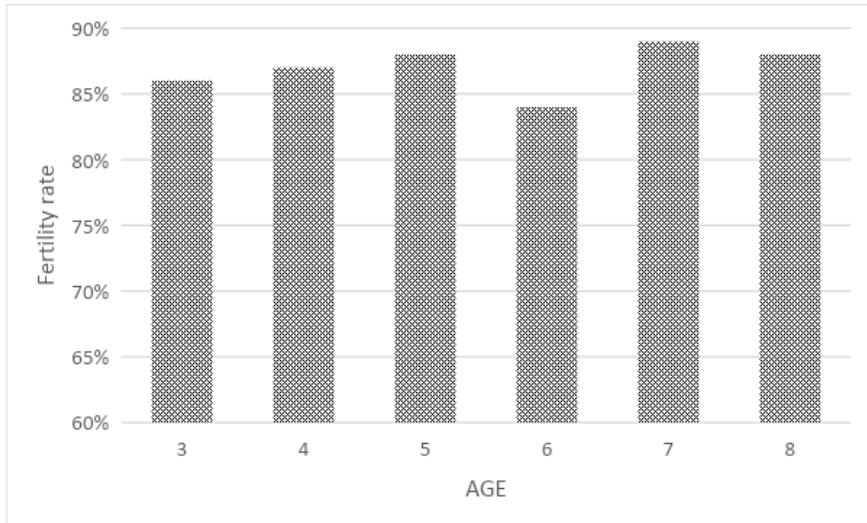


Figure S3. Graph showing fertility rate (%) divided by age, in the studied Lebanese Awassi population (n=165).