		Ge	Genotype I								G	enotyp	pe II												Genot	ype
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15	16	17	18	19	20	21	22	23	24	Γ
Genotype I	1. MG181946 KAS25-LOK-TR2017		100.0	100.0	97.0	97.7	99.0	99.0	98.7	98.7	99.3	99.3	99.3	99.3	99.3	99.7	99.7	99.7	99.3	99.3	99.0	98.7	98.7	98.7	91.7	4
	2. MG181944 KAS21-LOK-TR2017	100.0		100.0	97.0	97.7	99.0	99.0	98.7	98.7	99.3	99.3	99.3	99.3	99.3	99.7	99.7	99.7	99.3	99.3	99.0	98.7	98.7	98.7	91.7	1
	3. MG181945 KAS24-LOK-TR2017	100.0	100.0		97.0	97.7	99.0	99.0	98.7	98.7	99.3	99.3	99.3	99.3	99.3	99.7	99.7	99.7	99.3	99.3	99.0	98.7	98.7	98.7	91.7	4
		-	97.0	97.0	1	95.3	96.7	96.7	96.3	96.3	97.7	97.7	97.7	97.7	97.0	97.3	97.3	97.3	97.7	97.7	96.7	96.3	96.3	96.3	89.3	4
	5. GQ375280 YL	97.7	97.7	97.7	95.3		98.7	98.7	98.7	98.7	97.7	97.7	97.7		97.7	98.0	98.0			97.7	98.7	98.7	98.7	98.7	90.3	4
	6. KX192363 ANK-Br-TR2015	99.0	99.0	99.0	96.7	98.7		100.0	99.7	99.7	99.0	99.0	99.0	99.0	99.0	99.3	99.3	99.3	99.0	99.0	100.0	99.7	99.7	99.7	91.3	4
		-	99.0	99.0	96.7	98.7	100.0		99.7		_	99.0	99.0	99.0	99.0		_	99.3	99.0	99.0	100.0	99.7	99.7	99.7	91.3	4
		-	98.7	98.7	96.3	98.7	99.7	99.7		100.0	98.7	98.7	98.7	98.7	98.7	_	99.0	99.0	98.7	98.7	99.7	100.0	100.0	100.0	91.7	4
	9. GQ246866 strain TG-1	98.7	98.7	98.7	96.3	10000	99.7	99.7	100.0		98.7	98.7	98.7	98.7	98.7	99.0	99.0	10000	98.7	98.7	99.7	-	100.0	100.0	91.7	4
Genotype II	10. GQ246864 strain Kc-12	99.3	99.3	99.3	97.7	97.7	99.0	99.0	98.7	98.7		100.0			99.3	99.7	99.7	99.7	100.0	100.0	99.0	98.7	98.7	98.7	91.7	4
- 11			99.3	99.3	97.7	97.7	99.0	99.0	98.7	98.7	100.0				1000		2211	99.7		-	-			98.7	91.7	1
			99.3	99.3	97.7	97.7	99.0	99.0	98.7	98.7	100.0	100.0	_	100.0	_		-	99.7	_	_	-		-	98.7	91.7	4
		99.3	99.3	99.3	97.7	97.7	99.0	99.0	98.7	98.7	100.0	100.0	100.0		99.3		99.7	99.7	_	-		10000	98.7	98.7	91.7	4
			99.3	-	97.0	1	99.0	99.0	98.7	-	99.3		99.3	99.3		99.7	1220		-				10000	98.7	91.7	4
		-	-	-	-	98.0	99.3	99.3	99.0	99.0	99.7	99.7	99.7		99.7		100.0	100.0	_			_	-	99.0	92.0	4
		-	99.7	-	97.3	98.0	99.3	99.3	99.0	99.0	99.7	99.7	99.7	-	99.7	100.0		100.0	-	20000		2000		99.0	92.0	4
		10000	99.7	-	97.3	98.0	99.3	99.3	99.0	99.0	99.7	99.7	99.7	-	99.7	100.0	_	_	99.7			-		99.0	92.0	1
			99.3	-	2000	97.7	99.0	99.0	98.7	98.7	100.0	100.0	-	100.0	99.3	-	99.7	99.7		100.0				98.7	91.7	4
	The state of the s	-	99.3	99.3	97.7	97.7	99.0	99.0	98.7	98.7	100.0	100.0	100.0	100.0	99.3	99.7	99.7	99.7	100.0		99.0		-	98.7	91.7	4
		99.0	99.0	99.0	96.7	98.7	100.0	100.0	99.7	99.7	99.0	99.0	99.0	99.0	99.0	99.3	99.3	99.3	99.0	99.0		99.7	10000	99.7	91.3	4
	21. TR2011-TG-VS29 (MK543549)	98.7	98.7	98.7	96.3	98.7	99.7	99.7	100.0	100.0	98.7	98.7	98.7	98.7	98.7	99.0	99.0		98.7	98.7	99.7		100.0	100.0		4
		98.7	98.7	98.7	96.3	98.7	99.7	99.7	100.0	100.0	98.7	98.7	98.7	98.7	98.7	99.0	99.0	99.0	98.7	98.7	99.7	100.0		100.0	91.7	1
		98.7	98.7	98.7	96.3	98.7	99.7	99.7	100.0	100.0	98.7	98.7	98.7	98.7	98.7	99.0	99.0	99.0	98.7	98.7	99.7	100.0	100.0		91.7	4
Genotype III	24. MH605274 (9870MILK-TR2018)	91.7	91.7	91.7	89.3	90.3	91.3	91.3	91.7	91.7	91.7	91.7	91.7	91.7	91.7	92.0	92.0	92.0	91.7	91.7	91.3	91.7	91.7	91.7		1
																										Ī

Figure 3. *Nucleotid identity (%) of gB gene sequences of some Turkish BoHV-4.*