maternal chromosome 1										
maternar em omosome 1	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	2	7	16	25	20	17	5	cm sq	WI.	1 value
Pois	3.2	10.7	18.0	20.2	17.0	11.4	11.4	11.8	6	0.07
trunc Pois	3.1	10.7	18.1	20.2	17.0	11.4	11.4	11.7	6	0.07
free	2.0	7.1	16.0	24.4	23.5	13.8	5.3	2.0	0	0.07
trunc free	2.0	7.1	15.9	24.4	23.5	13.8	5.3	2.0		
gamma	1.0	6.2	16.8	24.2	22.8	14.1	7.0	3.7	5	0.59
gamma	1.0	0.2	10.0	27.2	22.0	17.1	7.0	3.7	3	0.57
maternal chromosome 2										
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	0	9	20	28	18	12	5	cm sq	W.	1 value
Pois	3.7	11.9	19.1	20.5	16.5	10.6	9.9	10.5	6	0.11
trunc Pois	3.6	12.0	19.2	20.5	16.4	10.5	9.8	10.3	6	0.11
free	1.5	8.2	19.4	25.7	21.0	11.1	5.0	2.3	0	0.11
trunc free	1.5	8.2	19.5	25.7	21.0	11.1	5.0	2.3		
gamma	1.6	8.2	18.9	25.2	20.8	11.9	5.3	2.4	5	0.78
gamma	1.0	0.2	10.7	23.2	20.0	11.7	3.3	2.4	3	0.76
maternal chromosome 3										
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	1	15	28	28	14	4	2	cm sq	-	1 varae
Pois	6.6	17.3	22.9	20.1	13.3	7.0	4.8	12.7	6	0.05
trunc Pois	6.3	17.7	23.1	20.2	13.2	6.9	4.6	12.1	6	0.06
free	3.0	14.0	26.8	26.7	15.1	5.0	1.4	2.8	0	0.00
trunc free	3.0	14.1	26.8	26.7	15.0	5.0	1.4	2.7		
gamma	3.4	14.6	25.1	25.6	15.7	5.7	1.9	3.4	5	0.64
gamma	3.4	14.0	23.1	23.0	13.7	3.7	1.7	5.4	3	0.04
maternal chromosome 4										
material en omosome 4	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	1	18	28	13	14	6	0	cm sq	41	1 value
Pois	6.6	16.5	20.6	17.1	10.6	5.3	3.3	13.1	5	0.02
trunc Pois	6.3	17.0	20.9	17.1	10.5	5.1	3.1	12.4	5	0.02
free	4.2	15.4	23.2	20.1	11.8	4.5	0.8	8.1	3	0.03
trunc free	4.2	15.4	23.2	20.1	11.8	4.5	0.8	8.1		
gamma	3.2	13.4	23.6	22.5	12.2	4.4	1.2	10.4	4	0.03
gamma	3.2	13.0	23.0	22.3	12.2	4.4	1.2	10.4	4	0.03
maternal chromosome 5										
maternar em omosome 5	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	5	15	26	21	8	5	0	cm-sq	uı	1 -value
Pois	7.7	18.1	21.1	16.4	9.6	4.5	2.6	6.8	5	0.24
trunc Pois	7.7	18.6	21.1	16.5	9.5	4.4	2.5	6.4	5	0.24
free	4.8	16.4	24.5	20.3	10.3	3.2	0.6	2.4	3	0.27
trunc free	4.6	16.4	24.6	20.5	10.3	3.1	0.5	2.4		
gamma	3.9	14.3	25.1	21.6	11.0	3.4	0.7	2.4	4	0.62
gamma	3.9	14.3	23.1	21.0	11.0	3.4	0.7	2.1	4	0.02

maternal chromosome 6										
maternar em omosome o	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	8	15	21	18	10	6	2	511 5 <b>q</b>		1 / 11/11/11
Pois	7.1	17.2	20.8	16.8	10.2	4.9	3.0	1.0	5	0.96
trunc Pois	6.6	17.5	21.0	16.9	10.2	4.9	2.9	1.3	5	0.94
free	7.1	16.8	20.1	17.5	10.9	5.0	2.5	0.7		0.7.
trunc free	7.0	16.9	20.1	17.5	10.9	5.0	2.5	0.8		
gamma	3.6	14.4	23.9	22.0	11.8	3.5	0.7	10.8	4	0.03
g	3.0	1	23.5	22.0	11.0	5.5	0.7	10.0		0.00
maternal chromosome 7										
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	10	13	30	25	11	3	0	- · · · · · ·		
Pois	9.7	21.8	24.5	18.4	10.4	4.7	2.5	10.3	5	0.07
trunc Pois	9.0	22.4	24.9	18.5	10.3	4.6	2.4	10.4	5	0.06
free	9.9	14.8	26.8	25.5	12.4	2.5	0.1	0.9		0.00
trunc free	7.7	18.4	25.9	25.1	12.3	2.5	0.1	3.2		
gamma	4.5	19.5	31.5	24.2	10.0	2.1	0.2	9.7	4	0.05
B	7.3	17.5	31.3	21.2	10.0	2.1	0.2	7.1	-	0.03
maternal chromosome 8										
maternal em omosome o	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	6	23	31	21	5	4	2	cm-sq	uı	1 -value
Pois	10.4	22.6	24.7	18.0	9.8	4.3	2.2	6.4	5	0.27
trunc Pois	9.7	23.4	25.1	18.0	9.6	4.1	2.1	5.5	5	0.27
free	6.7	23.4	30.5	19.2	7.1	2.9	2.3	1.3	3	0.30
trunc free	6.7	23.3	30.5	19.2	7.1	2.9	2.3	1.3		
	6.4	21.5	28.9	21.4	9.8	3.2	0.7	5.1	4	0.28
gamma	0.4	21.3	20.9	21.4	9.0	3.2	0.7	3.1	4	0.20
maternal chromosome 9										
maternal chromosome y	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	8	30	32	18	2	2	0	cm-sq	uı	1 -value
Pois	15.1	27.3	24.6	14.8	6.7	2.4	1.0	10.8	5	0.05
trunc Pois	13.1	29.2	25.2	14.5	6.2	2.1	0.8	9.0	5	0.03
free	9.3	29.2	32.8	15.6	3.6	1.0	0.8	2.4	3	0.11
trunc free	9.3	29.3	32.9	15.7	3.6	1.0	0.2	2.4		
gamma	9.3	27.8	31.2	17.8	5.0	0.9	0.3	3.6	4	0.46
gamma	9.1	21.0	31.2	17.0	3.0	0.9	0.1	3.0	4	0.40
maternal chromosome 10										
maternal em omosome 10	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	9	23	27	22	9	2	0	cm-sq	uı	1 -value
Pois	11.8	24.2	24.9	17.0	8.8	3.6	1.7	4.8	5	0.44
trunc Pois	10.9	25.2	25.4	17.0	8.5	3.4	1.6	4.3	5	0.44
free	9.1	22.6	27.6	21.6	9.2	1.8	0.1	0.2	3	0.51
trunc free	9.2	22.6	27.6	21.6	9.2	1.8	0.1	0.2		
	6.8	23.7	30.6	21.0	7.8	1.7	0.1	1.7	4	0.79
gamma	0.8	23.1	50.0	21.2	1.0	1./	0.3	1./	4	0.79
•	1								1	

0	1	2	3	4	5	5+	chi-sq	df	P-value
9	38	22	16	5	2	0	-		
16.2	28.1	24.4	14.2	6.2	2.1	0.8	8.2	5	0.15
									0.33
9.5	28.1	32.3	17.0	4.4	0.6	0.1	10.8	4	0.03
0	1	2	3	4	5	5+	chi-sq	df	P-value
2	24	28	17	8	1	0	_		
9.8	20.6	21.6	15.1	7.9	3.3	1.6	12.1	4	0.02
9.2	21.6	22.0	15.0	7.6	3.1	1.5	10.6	4	0.03
5.2	19.3	27.9	19.6	6.8	1.1	0.1	3.7		
5.2	19.3	28.0	19.6	6.8	1.1	0.1	3.7		
5.2	19.3	28.2	19.3	6.6	1.2	0.1	3.8	3	0.28
0	1	2	3	4	5	5+	chi-sq	df	P-value
23	37	28	3	1	0	0			
29.1	33.5	19.3	7.4	2.1	0.5	0.1	9.3	3	0.03
25.2	39.4	19.3	6.3	1.5	0.3	0.1	6.4	3	0.09
22.9	40.1	24.4	3.9	0.7	0.1	0.0	1.0		
20.7	42.2	24.6	3.7	0.7	0.1	0.0	1.6		
21.5	40.0	25.0	5.2	0.3	0.0	0.0	2.9	2	0.23
0	1	2	3	4	5	5+	chi-sa	df	P-value
				1					
				3.9			7.6	3	0.06
									0.09
								_	
					0.0			2	0.32
0	1	2	3	4	5	5+	chi-sq	df	P-value
20	34	31	7	0	0	0	-		
25.8	32.8	20.9	8.8	2.8	0.7	0.2	10.4	3	0.02
22.6	37.6	21.1	7.9	2.2	0.5	0.1	8.2	3	0.04
19.9	35.3	28.9	7.9	0.0	0.0	0.0	0.3		
17.9	38.4	28.1	7.6	0.0	0.0	0.0	1.1		
17.2	38.1	27.8	8.1	0.7	0.0	0.0	2.2	2	0.34
	9 16.2 14.8 12.5 12.4 9.5    9 9.8 9.2 5.2 5.2 5.2 5.2 5.2 5.2 12.5 12.5 12.	9 38  16.2 28.1  14.8 30.3  12.5 30.6  12.4 30.6  9.5 28.1  0 1  2 24  9.8 20.6  9.2 21.6  5.2 19.3  5.2 19.3  5.2 19.3  5.2 19.3  5.2 19.3  5.2 19.3  5.2 19.3  5.2 19.3  123 37  29.1 33.5  25.2 39.4  22.9 40.1  20.7 42.2  21.5 40.0  0 1  18 30  21.9 31.4  19.3 34.8  18.0 30.6  16.0 34.3  13.6 35.2  0 1  20 34  25.8 32.8  25.8 32.8  22.6 37.6  19.9 35.3  17.9 38.4	9 38 22  16.2 28.1 24.4  14.8 30.3 25.0  12.5 30.6 27.6  12.4 30.6 27.7  9.5 28.1 32.3  0 1 2  2 24 28  9.8 20.6 21.6  9.2 21.6 22.0  5.2 19.3 27.9  5.2 19.3 28.0  5.2 19.3 28.0  5.2 19.3 28.2  0 1 2  23 37 28  29.1 33.5 19.3  25.2 39.4 19.3  22.9 40.1 24.4  20.7 42.2 24.6  21.5 40.0 25.0  0 1 2  18 30 31  21.9 31.4 22.6  19.3 34.8 23.1  18.0 30.6 30.4  16.0 34.3 28.8  13.6 35.2 31.0  0 1 2  20 34 31  21.9 31.4 22.6  19.3 34.8 23.1  18.0 30.6 30.4  16.0 34.3 28.8  13.6 35.2 31.0	9         38         22         16           16.2         28.1         24.4         14.2           14.8         30.3         25.0         13.7           12.5         30.6         27.6         13.9           12.4         30.6         27.7         13.9           9.5         28.1         32.3         17.0           0         1         2         3           2         24         28         17           9.8         20.6         21.6         15.1           9.2         21.6         22.0         15.0           5.2         19.3         27.9         19.6           5.2         19.3         28.0         19.6           5.2         19.3         28.0         19.6           5.2         19.3         28.2         19.3           29.1         33.5         19.3         7.4           25.2         39.4         19.3         6.3           29.1         33.5         19.3         7.4           25.2         39.4         19.3         6.3           29.9         40.1         24.4         3.9           20.7         42.2	9         38         22         16         5           16.2         28.1         24.4         14.2         6.2           14.8         30.3         25.0         13.7         5.7           12.5         30.6         27.6         13.9         5.7           12.4         30.6         27.7         13.9         5.7           9.5         28.1         32.3         17.0         4.4           0         1         2         3         4           2         24         28         17         8           9.8         20.6         21.6         15.1         7.9           9.2         21.6         22.0         15.0         7.6           5.2         19.3         27.9         19.6         6.8           5.2         19.3         28.0         19.6         6.8           5.2         19.3         28.2         19.3         6.6           5.2         19.3         28.2         19.3         6.6           5.2         19.3         35.2         3         1           29.1         33.5         19.3         7.4         2.1           25.2         39.	9         38         22         16         5         2           16.2         28.1         24.4         14.2         6.2         2.1           14.8         30.3         25.0         13.7         5.7         1.9           12.5         30.6         27.6         13.9         5.7         1.5           12.4         30.6         27.7         13.9         5.7         1.5           9.5         28.1         32.3         17.0         4.4         0.6           0         1         2         3         4         5           2         24         28         17         8         1           9.8         20.6         21.6         15.1         7.9         3.3           9.2         21.6         22.0         15.0         7.6         3.1           5.2         19.3         27.9         19.6         6.8         1.1           5.2         19.3         28.0         19.6         6.8         1.1           5.2         19.3         28.2         19.3         6.6         1.2           0         1         2         3         4         5           2	9   38   22   16   5   2   0	9   38   22   16   5   2   0   16.2   28.1   24.4   14.2   6.2   2.1   0.8   8.2   14.8   30.3   25.0   13.7   5.7   1.9   0.7   5.7   12.5   30.6   27.6   13.9   5.7   1.5   0.2   4.7   12.4   30.6   27.7   13.9   5.7   1.5   0.2   4.7   9.5   28.1   32.3   17.0   4.4   0.6   0.1   10.8	9   38   22   16   5   2   0

4 11 17										
maternal chromosome 16	0	1	2	3	4	5	5+	ah! aa	df	P-value
Ol		37	28					chi-sq	aı	P-value
Obs Pois	10			12.9	2	1.7	0	0.1	4	0.06
	18.4	29.6	23.8	12.8	5.1		0.6	9.1	4	0.06
trunc Pois	16.8	32.4	24.3	12.1	4.5	1.4	0.4	6.0	4	0.20
free	11.9	32.7	31.6	12.7	2.5	0.6	0.1	1.6		
trunc free	11.8	32.7	31.6	12.7	2.5	0.6	0.1	1.6	2	0.44
gamma	12.0	31.5	30.2	14.3	3.5	0.4	0.0	2.9	3	0.41
maternal chromosome 17										
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	12	35	36	8	1	0	0	-		
Pois	21.2	31.1	22.8	11.2	4.1	1.2	0.4	16.8	3	0.00
trunc Pois	19.2	34.8	23.1	10.3	3.4	0.9	0.2	13.2	3	0.00
free	13.3	35.5	31.9	10.4	0.7	0.1	0.0	1.3		
trunc free	13.3	35.6	31.9	10.3	0.7	0.1	0.0	1.2		
gamma	14.0	34.7	30.9	10.5	1.8	0.1	0.0	2.2	2	0.33
maternal chromosome 18	0	1	2	2	4	-	5+	-1.2	df	P-value
Obs	11	31	28	7	3	5	0	chi-sq	aı	P-value
Pois	17.9	26.8	20.1	10.0	3.8	1.1		9.0	4	0.06
trunc Pois	16.1	29.7	20.1	9.4	3.8	0.9	0.4	6.2	4	0.08
free	12.6	30.6	25.3	9.4	2.1	0.9	0.0	1.6	4	0.18
trunc free	12.6 10.7	30.7	25.4	9.2	2.1	0.2	0.0	1.6	2	0.52
gamma	10.7	29.4	26.8	10.8	2.0	0.2	0.0	2.2	3	0.53
maternal chromosome 19	0	1	2	2	4	_	<i>-</i> .	-1-2	df	P-value
Obs	26	40	23	3	0	5	5+ 0	chi-sq	aı	P-value
Pois		33.8		6.0				7.7	2	0.05
	32.8		17.5		1.6	0.3	0.1	7.7	3	0.05
trunc Pois	28.1	41.2	16.9	4.6	0.9	0.2	0.0	4.1	3	0.25
free	25.9	40.4	22.5	3.1	0.0	0.0	0.0	0.0		
trunc free	24.3	42.9	21.6	3.1	0.0	0.0	0.0	0.5	-	0.50
gamma	23.2	42.6	22.8	3.4	0.1	0.0	0.0	0.6	2	0.73
maternal chromosome 20										
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	22	43	22	5	0	0	0	-		
Pois	30.4	33.7	18.7	6.9	1.9	0.4	0.1	8.4	3	0.04
trunc Pois	26.6	40.3	18.1	5.4	1.2	0.2	0.0	3.3	3	0.35
free	23.2	40.9	23.1	4.8	0.1	0.0	0.0	0.3		
trunc free	22.9	41.3	23.1	4.7	0.0	0.0	0.0	0.2		
gamma	22.6	38.6	23.7	6.1	0.9	0.0	0.0	1.8	2	0.41

maternal chromosome 21										
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	40	35	5	0	0	0	0			
Pois	45.6	25.6	7.2	1.4	0.2	0.0	0.0	6.3	2	0.04
trunc Pois	36.0	39.7	4.0	0.3	0.0	0.0	0.0	1.5	2	0.46
free	40.0	35.1	4.9	0.0	0.0	0.0	0.0	0.0		0.10
trunc free	35.6	39.9	4.4	0.1	0.0	0.0	0.0	1.3		
gamma	38.8	35.5	5.6	0.1	0.0	0.0	0.0	0.3	1	0.61
gamma	30.0	33.3	5.0	0.1	0.0	0.0	0.0	0.5	1	0.01
maternal chromosome 22										
maternal chromosome 22	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	49	38	5	0	0	0	0	cm-sq	uı	1 -value
Pois	54.6	28.5	7.4	1.3	0.2	0.0	0.0	6.0	2	0.05
trunc Pois	42.1	45.8	3.9	0.2	0.2	0.0	0.0	3.0	2	0.03
free		38.1		0.2		0.0	0.0	0.0		0.22
	49.0		4.9		0.0					
trunc free	41.8	45.9	4.2	0.1	0.0	0.0	0.0	2.8	1	0.73
gamma	49.2	37.9	4.8	0.1	0.0	0.0	0.0	0.1	1	0.72
4 11										
maternal chromosome X				2		_			7.0	
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	11	27	33	17	3	1	0			
Pois	16.0	28.0	24.5	14.3	6.2	2.2	0.8	8.1	4	0.09
trunc Pois	14.5	29.9	25.1	14.0	5.9	2.0	0.7	6.8	4	0.15
free	10.9	28.2	32.1	16.3	3.8	0.6	0.1	0.4		
trunc free	10.2	29.2	32.0	16.2	3.8	0.6	0.1	0.6		
gamma	9.1	28.7	32.5	17.5	3.8	0.4	0.0	1.5	3	0.69
					<del></del>		-			

paternal chromosome 1										
<b>partition</b> 0.11 0.11 0.11 0.11 0.11 0.11 0.11 0.1	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	8	30	30	19	5	0	0	<b>1</b>		
Pois	15.0	27.2	24.7	14.9	6.8	2.5	1.0	9.7	4	0.05
trunc Pois	13.8	29.0	25.2	14.6	6.3	2.2	0.8	8.0	4	0.09
free	9.0	27.3	32.5	18.6	4.5	0.1	0.0	0.7	1	0.40
trunc free	8.9	27.4	32.7	18.6	4.4	0.0	0.0	0.7	1	0.41
gamma	8.6	26.7	32.3	18.3	5.2	1.0	0.0	1.7	3	0.41
gamma	0.0	20.7	32.3	10.5	3.2	1.0	0.1	1.7		0.04
paternal chromosome 2										
<b>P</b>	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	7	27	31	20	7	0	0	1		
Pois	13.4	25.8	24.9	15.9	7.7	3.0	1.3	10.0	4	0.04
trunc Pois	12.4	27.3	25.4	15.7	7.3	2.7	1.1	8.6	4	0.07
free	7.6	25.2	32.9	20.7	5.4	0.2	0.0	0.9	1	0.34
trunc free	7.6	25.2	33.0	20.6	5.4	0.2	0.0	0.9	1	0.34
gamma	8.1	25.8	31.3	19.4	6.3	1.0	0.1	1.5	3	0.69
D	0.1	23.0	31.3	17.7	5.5	1.0	0.1	1.5	5	0.07
paternal chromosome 3										
paternar em omosome s	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	7.0	37.0	32.0	15.0	1.0	0.0	0.0	cm-sq	uı	1 - varue
Pois	18.0	29.4	23.9	13.0	5.3	1.7	0.6	17.5	3	0.00
trunc Pois	16.7	32.3	24.3	12.2	4.6	1.4	0.4	13.9	3	0.00
free	10.7	32.3	34.3	13.8	1.3	0.0	0.0	2.2	1	0.14
trunc free	10.4	32.2	34.3	13.8	1.3	0.0	0.0	2.2	1	0.14
	10.4	30.3	32.8	15.5	2.9	0.0	0.0	4.0	2	0.14
gamma	10.2	30.3	32.0	13.3	2.9	0.2	0.0	4.0		0.13
paternal chromosome 4										
paternal chromosome 4	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	10	39	32	10	1	0	0	cm-sq	uı	r-value
Pois	20.8	30.9	23.0	11.4	4.3	1.3	0.4	15.5	3	0.00
trunc Pois	19.0	34.6	23.3	10.4	3.5	0.9	0.4	11.1	3	0.00
	12.9	35.0	32.3	11.0	0.8	0.9	0.3	11.1	1	0.01
free			32.3					1.2		
trunc free	12.9	35.1		10.9	0.8	0.0	0.0		1	0.28
gamma	13.1	34.3	30.0	12.4	2.1	0.1	0.0	2.6	2	0.27
mataumal abusama asuma 5										
paternal chromosome 5	0	1	2	3	4	5	5+	a <b>la:</b> a a	df	P-value
Ol-	0	1			4			chi-sq	aı	P-value
Obs	18	38	30	6	0	0	0	11.0	2	0.01
Pois	26.1	32.9	20.7	8.7	2.7	0.7	0.2	11.9	3	0.01
trunc Pois	23.1	38.0	20.7	7.5	2.1	0.4	0.1	8.2	3	0.04
free	17.9	39.1	28.4	6.5	0.0	0.0	0.0	0.2	1	0.65
trunc free	17.4	39.7	28.5	6.3	0.0	0.0	0.0	0.2	1	0.64
gamma	19.2	37.5	26.2	7.9	1.1	0.0	0.0	2.2	2	0.33

0	1	2	3	4	5	5+	chi-sq	df	P-value
22	36	30	4	0	0	0	541 S4		
							11.8	3	0.01
									0.03
									0.35
									0.20
									0.36
17.0	40.5	20.0	3.7	0.2	0.0	0.0	2.1		0.50
0	1	2	3	4	5	5+	chi-sa	df	P-value
							ciii sq	ui.	1 value
					-	-	4 1	3	0.25
									0.78
									0.78
									0.51
									0.32
21.0	40.3	24.1	0.0	0.0	0.0	0.0	2.2	2	0.34
0	1	2	2	1	5	5.	chi sa	4t	P-value
							cm-sq	uı	r-value
	_						12.7	2	0.01
									0.01
									0.13
									0.63
24.8	42.0	21.2	3.8	0.1	0.0	0.0	1.5	2	0.48
0	1	2	2	4	-	<b>-</b> .	1 •	10	D 1
							cm-sq	aı	P-value
					_	-	11.6	2	0.01
									0.01
									0.09
									0.83
									0.79
21.7	40.4	23.3	6.0	0.7	0.0	0.0	1.9	2	0.39
								7.0	D 1
	1	•	2	4	_		1 .		P-value
0	1	2	3	4	5	5+	chi-sq	df	
17	39	28	8	0	0	0	-		0.00
17 25.2	<b>39</b> 32.6	<b>28</b> 21.1	<b>8</b> 9.1	<b>0</b> 2.9	0.8	0.2	10.2	3	0.02
25.2 22.5	39 32.6 37.5	28 21.1 21.2	9.1 8.0	2.9 2.3	0 0.8 0.5	0.2 0.1	10.2	3	0.09
25.2 22.5 17.4	39 32.6 37.5 38.1	28 21.1 21.2 28.7	8 9.1 8.0 7.7	2.9 2.3 0.0	0.8 0.5 0.0	0 0.2 0.1 0.0	10.2 6.5 0.1	3 3 1	0.09 0.76
25.2 22.5 17.4 17.2	39 32.6 37.5 38.1 38.4	28 21.1 21.2 28.7 28.8	8 9.1 8.0 7.7 7.6	2.9 2.3 0.0 0.0	0.8 0.5 0.0 0.0	0.2 0.1 0.0 0.0	10.2 6.5 0.1 0.1	3 3 1 1	0.09 0.76 0.80
25.2 22.5 17.4	39 32.6 37.5 38.1	28 21.1 21.2 28.7	8 9.1 8.0 7.7	2.9 2.3 0.0	0.8 0.5 0.0	0 0.2 0.1 0.0	10.2 6.5 0.1	3 3 1	0.09 0.76
25.2 22.5 17.4 17.2	39 32.6 37.5 38.1 38.4	28 21.1 21.2 28.7 28.8	8 9.1 8.0 7.7 7.6	2.9 2.3 0.0 0.0	0.8 0.5 0.0 0.0	0.2 0.1 0.0 0.0	10.2 6.5 0.1 0.1	3 3 1 1	0.09 0.76 0.80
25.2 22.5 17.4 17.2	39 32.6 37.5 38.1 38.4	28 21.1 21.2 28.7 28.8	8 9.1 8.0 7.7 7.6	2.9 2.3 0.0 0.0	0.8 0.5 0.0 0.0	0.2 0.1 0.0 0.0	10.2 6.5 0.1 0.1	3 3 1 1	0.09 0.76 0.80
25.2 22.5 17.4 17.2	39 32.6 37.5 38.1 38.4	28 21.1 21.2 28.7 28.8	8 9.1 8.0 7.7 7.6	2.9 2.3 0.0 0.0	0.8 0.5 0.0 0.0	0.2 0.1 0.0 0.0	10.2 6.5 0.1 0.1	3 3 1 1	0.09 0.76 0.80
25.2 22.5 17.4 17.2	39 32.6 37.5 38.1 38.4	28 21.1 21.2 28.7 28.8	8 9.1 8.0 7.7 7.6	2.9 2.3 0.0 0.0	0.8 0.5 0.0 0.0	0.2 0.1 0.0 0.0	10.2 6.5 0.1 0.1	3 3 1 1	0.09 0.76 0.80
25.2 22.5 17.4 17.2	39 32.6 37.5 38.1 38.4	28 21.1 21.2 28.7 28.8	8 9.1 8.0 7.7 7.6	2.9 2.3 0.0 0.0	0.8 0.5 0.0 0.0	0.2 0.1 0.0 0.0	10.2 6.5 0.1 0.1	3 3 1 1	0.09 0.76 0.80
25.2 22.5 17.4 17.2	39 32.6 37.5 38.1 38.4	28 21.1 21.2 28.7 28.8	8 9.1 8.0 7.7 7.6	2.9 2.3 0.0 0.0	0.8 0.5 0.0 0.0	0.2 0.1 0.0 0.0	10.2 6.5 0.1 0.1	3 3 1 1	0.09 0.76 0.80
	22 28.4 24.7 21.9 19.7 19.0  0 23 28.8 24.9 24.5 21.0  0 23 33.1 29.2 24.2 24.2 24.8  0 20 29.1 25.7 20.4 20.3 21.7	28.4 33.4 24.7 39.1 21.9 38.7 19.7 41.3 19.0 40.3 19.0 40.3 23 41 28.8 33.4 24.9 39.2 24.5 38.6 21.0 40.3 24.5 38.6 21.0 40.3 29.2 41.8 24.2 44.0 24.2 44.1 24.8 42.0 24.2 44.1 24.8 42.0 25.7 39.8 20.4 41.7 20.3 41.9	28.4       33.4       19.6         24.7       39.1       19.6         21.9       38.7       26.5         19.7       41.3       26.2         19.0       40.3       26.8         0       1       2         23       41       19         28.8       33.4       19.4         24.9       39.2       19.5         24.5       38.6       20.5         24.5       38.6       20.5         21.0       40.3       24.1         0       1       2         23       46       21         33.1       33.8       17.3         29.2       41.8       16.0         24.2       44.0       21.9         24.2       44.1       21.8         24.8       42.0       21.2         0       1       2         20       42       26         29.1       33.5       19.3         25.7       39.8       18.8         20.4       41.7       25.8         20.3       41.9       25.6	28.4       33.4       19.6       7.7         24.7       39.1       19.6       6.5         21.9       38.7       26.5       5.0         19.7       41.3       26.2       4.7         19.0       40.3       26.8       5.7         0       1       2       3         23       41       19       8         28.8       33.4       19.4       7.5         24.9       39.2       19.5       6.5         24.5       38.6       20.5       7.3         24.5       38.6       20.5       7.3         21.0       40.3       24.1       6.0         0       1       2       3         23       46       21       2         33.1       33.8       17.3       5.9         29.2       41.8       16.0       4.1         24.2       44.0       21.9       1.9         24.2       44.1       21.8       1.9         24.8       42.0       21.2       3.8         0       1       2       3         20       42       26       4         29.1	28.4       33.4       19.6       7.7       2.3         24.7       39.1       19.6       6.5       1.6         21.9       38.7       26.5       5.0       0.0         19.7       41.3       26.2       4.7       0.1         19.0       40.3       26.8       5.7       0.2         0       1       2       3       4         23       41       19       8       1         28.8       33.4       19.4       7.5       2.2         24.9       39.2       19.5       6.5       1.6         24.5       38.6       20.5       7.3       1.1         24.5       38.6       20.5       7.3       1.1         21.0       40.3       24.1       6.0       0.6         33.1       33.8       17.3       5.9       1.5         29.2       41.8       16.0       4.1       0.8         24.2       44.0       21.9       1.9       0.0         24.2       44.1       21.8       1.9       0.1         24.8       42.0       21.2       3.8       0.1         0       1       2 <t< td=""><td>28.4         33.4         19.6         7.7         2.3         0.5           24.7         39.1         19.6         6.5         1.6         0.3           21.9         38.7         26.5         5.0         0.0         0.0           19.7         41.3         26.2         4.7         0.1         0.0           19.0         40.3         26.8         5.7         0.2         0.0           19.0         40.3         26.8         5.7         0.2         0.0           28.8         33.4         19.4         7.5         2.2         0.5           24.9         39.2         19.5         6.5         1.6         0.3           24.5         38.6         20.5         7.3         1.1         0.0           24.5         38.6         20.5         7.3         1.1         0.1           21.0         40.3         24.1         6.0         0.6         0.0           33.1         33.8         17.3         5.9         1.5         0.3           29.2         41.8         16.0         4.1         0.8         0.1           24.2         44.0         21.9         1.9         0.0</td><td>28.4       33.4       19.6       7.7       2.3       0.5       0.1         24.7       39.1       19.6       6.5       1.6       0.3       0.1         21.9       38.7       26.5       5.0       0.0       0.0       0.0         19.7       41.3       26.2       4.7       0.1       0.0       0.0         19.0       40.3       26.8       5.7       0.2       0.0       0.0         19.0       40.3       26.8       5.7       0.2       0.0       0.0         28.8       33.4       19.4       7.5       2.2       0.5       0.1         24.9       39.2       19.5       6.5       1.6       0.3       0.1         24.5       38.6       20.5       7.3       1.1       0.0       0.0         24.5       38.6       20.5       7.3       1.1       0.1       0.0         24.0       40.3       24.1       6.0       0.6       0.0       0.0         21.0       40.3       24.1       6.0       0.6       0.0       0.0         24.2       44.0       21.9       1.9       0.0       0.0       0.0</td><td>28.4         33.4         19.6         7.7         2.3         0.5         0.1         11.8           24.7         39.1         19.6         6.5         1.6         0.3         0.1         9.1           21.9         38.7         26.5         5.0         0.0         0.0         0.0         0.9           19.7         41.3         26.2         4.7         0.1         0.0         0.0         1.6           19.0         40.3         26.8         5.7         0.2         0.0         0.0         2.1           0         1         2         3         4         5         5+         chi-sq           23         41         19         8         1         0         0           24.9         39.2         19.5         6.5         1.6         0.3         0.1         1.1           24.9         39.2         19.5         6.5         1.6         0.3         0.1         1.1           24.5         38.6         20.5         7.3         1.1         0.0         0.0         0.4           24.5         38.6         20.5         7.3         1.1         0.1         0.0         0.0</td><td>28.4 33.4 19.6 7.7 2.3 0.5 0.1 11.8 3 24.7 39.1 19.6 6.5 1.6 0.3 0.1 9.1 3 21.9 38.7 26.5 5.0 0.0 0.0 0.0 0.0 0.0 0.9 1 19.7 41.3 26.2 4.7 0.1 0.0 0.0 0.0 1.6 1 19.0 40.3 26.8 5.7 0.2 0.0 0.0 0.0 2.1 2   0 1 2 3 4 19 8 1 0 0 28.8 33.4 19.4 7.5 2.2 0.5 0.1 4.1 3 24.9 39.2 19.5 6.5 1.6 0.3 0.1 1.1 3 24.5 38.6 20.5 7.3 1.1 0.0 0.0 0.0 0.4 1 24.5 38.6 20.5 7.3 1.1 0.0 0.0 0.0 0.4 1 21.0 40.3 24.1 6.0 0.6 0.0 0.0 0.0 2.2 2   0 1 2 3 46 21 2 0 0 0 33.1 33.8 17.3 5.9 1.5 0.3 0.1 12.7 3 29.2 41.8 16.0 4.1 0.8 0.1 0.0 5.3 3 24.2 44.0 21.9 1.9 0.0 0.0 0.0 0.2 1 24.2 44.1 21.8 1.9 0.1 0.0 0.0 0.2 1 24.8 42.0 21.2 3.8 0.1 0.0 0.0 0.0 1.5 2  0 1 2 3 4 5 5+ chi-sq df 20 42 26 4 0 0 0 29.1 33.5 19.3 7.4 2.1 0.5 0.1 11.6 3 25.7 39.8 18.8 5.9 1.4 0.3 0.0 0.0 0.1 1 20.3 41.9 25.6 4.1 0.1 0.0 0.0 0.0 0.1 1</td></t<>	28.4         33.4         19.6         7.7         2.3         0.5           24.7         39.1         19.6         6.5         1.6         0.3           21.9         38.7         26.5         5.0         0.0         0.0           19.7         41.3         26.2         4.7         0.1         0.0           19.0         40.3         26.8         5.7         0.2         0.0           19.0         40.3         26.8         5.7         0.2         0.0           28.8         33.4         19.4         7.5         2.2         0.5           24.9         39.2         19.5         6.5         1.6         0.3           24.5         38.6         20.5         7.3         1.1         0.0           24.5         38.6         20.5         7.3         1.1         0.1           21.0         40.3         24.1         6.0         0.6         0.0           33.1         33.8         17.3         5.9         1.5         0.3           29.2         41.8         16.0         4.1         0.8         0.1           24.2         44.0         21.9         1.9         0.0	28.4       33.4       19.6       7.7       2.3       0.5       0.1         24.7       39.1       19.6       6.5       1.6       0.3       0.1         21.9       38.7       26.5       5.0       0.0       0.0       0.0         19.7       41.3       26.2       4.7       0.1       0.0       0.0         19.0       40.3       26.8       5.7       0.2       0.0       0.0         19.0       40.3       26.8       5.7       0.2       0.0       0.0         28.8       33.4       19.4       7.5       2.2       0.5       0.1         24.9       39.2       19.5       6.5       1.6       0.3       0.1         24.5       38.6       20.5       7.3       1.1       0.0       0.0         24.5       38.6       20.5       7.3       1.1       0.1       0.0         24.0       40.3       24.1       6.0       0.6       0.0       0.0         21.0       40.3       24.1       6.0       0.6       0.0       0.0         24.2       44.0       21.9       1.9       0.0       0.0       0.0	28.4         33.4         19.6         7.7         2.3         0.5         0.1         11.8           24.7         39.1         19.6         6.5         1.6         0.3         0.1         9.1           21.9         38.7         26.5         5.0         0.0         0.0         0.0         0.9           19.7         41.3         26.2         4.7         0.1         0.0         0.0         1.6           19.0         40.3         26.8         5.7         0.2         0.0         0.0         2.1           0         1         2         3         4         5         5+         chi-sq           23         41         19         8         1         0         0           24.9         39.2         19.5         6.5         1.6         0.3         0.1         1.1           24.9         39.2         19.5         6.5         1.6         0.3         0.1         1.1           24.5         38.6         20.5         7.3         1.1         0.0         0.0         0.4           24.5         38.6         20.5         7.3         1.1         0.1         0.0         0.0	28.4 33.4 19.6 7.7 2.3 0.5 0.1 11.8 3 24.7 39.1 19.6 6.5 1.6 0.3 0.1 9.1 3 21.9 38.7 26.5 5.0 0.0 0.0 0.0 0.0 0.0 0.9 1 19.7 41.3 26.2 4.7 0.1 0.0 0.0 0.0 1.6 1 19.0 40.3 26.8 5.7 0.2 0.0 0.0 0.0 2.1 2   0 1 2 3 4 19 8 1 0 0 28.8 33.4 19.4 7.5 2.2 0.5 0.1 4.1 3 24.9 39.2 19.5 6.5 1.6 0.3 0.1 1.1 3 24.5 38.6 20.5 7.3 1.1 0.0 0.0 0.0 0.4 1 24.5 38.6 20.5 7.3 1.1 0.0 0.0 0.0 0.4 1 21.0 40.3 24.1 6.0 0.6 0.0 0.0 0.0 2.2 2   0 1 2 3 46 21 2 0 0 0 33.1 33.8 17.3 5.9 1.5 0.3 0.1 12.7 3 29.2 41.8 16.0 4.1 0.8 0.1 0.0 5.3 3 24.2 44.0 21.9 1.9 0.0 0.0 0.0 0.2 1 24.2 44.1 21.8 1.9 0.1 0.0 0.0 0.2 1 24.8 42.0 21.2 3.8 0.1 0.0 0.0 0.0 1.5 2  0 1 2 3 4 5 5+ chi-sq df 20 42 26 4 0 0 0 29.1 33.5 19.3 7.4 2.1 0.5 0.1 11.6 3 25.7 39.8 18.8 5.9 1.4 0.3 0.0 0.0 0.1 1 20.3 41.9 25.6 4.1 0.1 0.0 0.0 0.0 0.1 1

paternal chromosome 11										
paternar em omosome 11	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	23	49	17	3	0	0	0	<b>1</b>		
Pois	33.8	33.8	16.9	5.6	1.4	0.3	0.1	13.2	3	0.00
trunc Pois	30.0	42.2	15.3	3.7	0.7	0.1	0.0	3.8	3	0.28
free	26.5	43.3	19.5	2.6	0.1	0.0	0.0	1.7	1	0.20
trunc free	26.4	43.4	19.6	2.6	0.1	0.0	0.0	1.6	1	0.20
gamma	25.9	40.1	20.7	4.7	0.6	0.0	0.0	4.2	2	0.13
gamma	23.7	70.1	20.7	т. /	0.0	0.0	0.0	7.2		0.13
paternal chromosome 12										
paternar em omosome 12	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	21	40	26	4	1	0	0	cm sq	41	1 value
Pois	28.4	33.4	19.6	7.7	2.3	0.5	0.1	8.4	3	0.04
trunc Pois	24.9	39.3	19.4	6.4	1.6	0.3	0.1	4.2	3	0.24
free	20.9	41.0	24.9	4.4	0.8	0.3	0.0	0.1	1	0.72
trunc free	20.4	41.5	24.8	4.4	0.8	0.1	0.0	0.1	1	0.72
	20.4	39.9	25.4	5.7	0.8	0.1	0.0	1.3	2	0.67
gamma	20.3	37.7	23.4	3.1	0.4	0.0	0.0	1.3		0.32
paternal chromosome 13										
paternar cin omosome 13	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	36	42	14	0	0	0	0	cm-sq	uı	r-value
Pois		32.7		3.2				7.0	2	0.02
trunc Pois	43.0		12.4		0.6	0.1	0.0	7.8	2	0.02
	35.6	44.4	10.2	1.6	0.2	0.0	0.0		2	0.19
free	35.9	42.2	13.9	0.0	0.0	0.0	0.0	0.0	1	0.95
trunc free	33.2	46.0	12.8	0.0	0.0	0.0	0.0	0.7	1	0.40
gamma	36.2	42.9	12.0	0.9	0.0	0.0	0.0	1.3	1	0.25
paternal chromosome 14						_	_			
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	27	46	19	0	0	0	0			
Pois	36.9	33.7	15.4	4.7	1.1	0.2	0.0	14.0	2	0.00
trunc Pois	32.2	43.2	13.4	2.8	0.4	0.1	0.0	6.6	2	0.04
free	27.3	45.5	19.1	0.0	0.0	0.0	0.0	0.0	1	0.89
trunc free	27.1	46.0	18.9	0.0	0.0	0.0	0.0	0.0	1	0.93
gamma	25.1	43.3	20.5	3.0	0.1	0.0	0.0	3.6	1	0.06
paternal chromosome 15				_						
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	36	44	12	0	0	0	0			
Pois	43.9	32.5	12.0	3.0	0.5	0.1	0.0	9.1	2	0.01
trunc Pois	36.7	44.7	9.2	1.3	0.1	0.0	0.0	2.3	2	0.32
free	36.0	44.0	12.0	0.0	0.0	0.0	0.0	0.0	1	0.88
trunc free	34.5	46.0	11.5	0.0	0.0	0.0	0.0	0.2	1	0.66
gamma	35.3	43.4	12.5	0.8	0.0	0.0	0.0	0.9	1	0.35

paternal chromosome 16	0	1	2	3	4	5	5+	a <b>l.:</b> a a	df	P-value
OL -			13	1				chi-sq	aı	P-value
Obs Pois	32	46		3.7	0	0	0	0.6	2	0.01
	40.7	33.2	13.5		0.7	0.1	0.0	9.6	2	0.01
trunc Pois	34.7	44.1	11.1	1.9	0.2	0.0	0.0	1.2	2	0.55
free	32.9	44.7	13.4	0.9	0.0	0.0	0.0	0.1	1	0.78
trunc free	32.7	45.0	13.2	1.0	0.1	0.0	0.0	0.0	1	0.84
gamma	30.7	45.7	15.0	0.6	0.0	0.0	0.0	0.6	1	0.45
paternal chromosome 17										
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	33	48	11	0	0	0	0	_		
Pois	43.0	32.7	12.4	3.2	0.6	0.1	0.0	13.5	2	0.00
trunc Pois	36.9	44.8	9.0	1.2	0.1	0.0	0.0	2.4	2	0.30
free	34.7	45.7	11.6	0.0	0.0	0.0	0.0	0.2	1	0.63
trunc free	34.5	46.0	11.5	0.0	0.0	0.0	0.0	0.2	1	0.66
gamma	33.3	44.6	13.5	0.6	0.0	0.0	0.0	1.3	1	0.26
paternal chromosome 18				2		_	_		10	
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	31	47	14	0	0	0	0			
Pois	40.7	33.2	13.5	3.7	0.7	0.1	0.0	12.7	2	0.00
trunc Pois	35.1	44.2	10.7	1.7	0.2	0.0	0.0	3.6	2	0.16
free	32.0	45.4	14.5	0.0	0.0	0.0	0.0	0.1	1	0.72
trunc free	31.7	46.0	14.3	0.0	0.0	0.0	0.0	0.1	1	0.81
gamma	30.8	44.4	15.4	1.3	0.0	0.0	0.0	1.6	1	0.21
paternal chromosome 19										
	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	40	36	16	0	0	0	0			
Pois	43.9	32.5	12.0	3.0	0.5	0.1	0.0	5.7	2	0.06
trunc Pois	35.1	44.2	10.7	1.7	0.2	0.0	0.0	6.8	2	0.03
free	40.0	36.3	15.8	0.0	0.0	0.0	0.0	0.0	1	0.94
trunc free	32.9	46.0	13.1	0.0	0.0	0.0	0.0	4.4	1	0.04
gamma	35.4	43.8	12.3	0.5	0.0	0.0	0.0	3.6	1	0.06
paternal chromosome 20										
paternar em omosome 20	0	1	2	3	4	5	5+	chi-sq	df	P-value
Obs	48	38	6	0	0	0	0	em sq	WI.	1 value
Pois	53.4	29.0	7.9	1.4	0.2	0.0	0.0	5.4	2	0.07
trunc Pois	41.3	45.7	4.7	0.3	0.2	0.0	0.0	3.4	2	0.07
free	48.0	38.1	5.9	0.0	0.0	0.0	0.0	0.0		0.21
trunc free	40.9	45.9	5.1	0.0	0.0	0.0	0.0	2.8		
gamma	46.4	38.5	6.8	0.1	0.0	0.0	0.0	0.5	1	0.49
Ramma	40.4	20.2	0.8	0.3	0.0	0.0	0.0	0.3	1	0.49

0	1	2	3	4	5	5+	chi-sq	df	P-value
58	34	0	0	0	0	0			
63.6	23.5	4.3	0.5	0.0	0.0	0.0	10.1	1	0.00
46.0	46.0	0.0	0.0	0.0	0.0	0.0	6.3	1	0.01
58.0	34.0	0.0	0.0	0.0	0.0	0.0	0.0	1	0.91
46.0	46.0	0.0	0.0	0.0	0.0	0.0	6.3	1	0.01
0	1	2	3	4	5	5+	chi-sq	df	P-value
69	22	1	0	0	0	0			
70.9	18.5	2.4	0.2	0.0	0.0	0.0	1.7	1	0.19
45.3	46.0	0.7	0.0	0.0	0.0	0.0	25.0	1	0.00
69.0	22.0	1.0	0.0	0.0	0.0	0.0	0.0	1	0.98
45.3	46.0	0.7	0.0	0.0	0.0	0.0	25.0	1	0.00
67.3	24.2	0.5	0.0	0.0	0.0	0.0	0.7	0	
0	1	2	3	4	5	5+	chi-sq	df	P-value
79	13	0	0	0	0	0			
79.9	11.3	0.8	0.0	0.0	0.0	0.0	1.1	1	0.29
46.0	46.0	0.0	0.0	0.0	0.0	0.0	47.4	1	0.00
79.0	13.0	0.0	0.0	0.0	0.0	0.0	0.0	1	0.93
46.0	46.0	0.0	0.0	0.0	0.0	0.0	47.4	1	0.00
	58 63.6 46.0 58.0 46.0 69 70.9 45.3 69.0 45.3 67.3	58         34           63.6         23.5           46.0         46.0           58.0         34.0           46.0         46.0           69         22           70.9         18.5           45.3         46.0           69.0         22.0           45.3         46.0           67.3         24.2           0         1           79         13           79.9         11.3           46.0         46.0           79.0         13.0	58         34         0           63.6         23.5         4.3           46.0         46.0         0.0           58.0         34.0         0.0           46.0         46.0         0.0           46.0         46.0         0.0           69         22         1           70.9         18.5         2.4           45.3         46.0         0.7           69.0         22.0         1.0           45.3         46.0         0.7           67.3         24.2         0.5           0         1         2           79         13         0           79.9         11.3         0.8           46.0         46.0         0.0           79.0         13.0         0.0	58         34         0         0           63.6         23.5         4.3         0.5           46.0         46.0         0.0         0.0           58.0         34.0         0.0         0.0           46.0         46.0         0.0         0.0           46.0         46.0         0.0         0.0           70.9         18.5         2.4         0.2           45.3         46.0         0.7         0.0           69.0         22.0         1.0         0.0           45.3         46.0         0.7         0.0           67.3         24.2         0.5         0.0           0         1         2         3           79         13         0         0           79.9         11.3         0.8         0.0           46.0         46.0         0.0         0.0           79.0         13.0         0.0         0.0	58         34         0         0         0           63.6         23.5         4.3         0.5         0.0           46.0         46.0         0.0         0.0         0.0           58.0         34.0         0.0         0.0         0.0           46.0         46.0         0.0         0.0         0.0           46.0         46.0         0.0         0.0         0.0           70.9         18.5         2.4         0.2         0.0           45.3         46.0         0.7         0.0         0.0           45.3         46.0         0.7         0.0         0.0           67.3         24.2         0.5         0.0         0.0           67.3         24.2         0.5         0.0         0.0           79.9         11.3         0.8         0.0         0.0           46.0         46.0         0.0         0.0         0.0           79.0         13.0         0.0         0.0         0.0	58         34         0         0         0         0           63.6         23.5         4.3         0.5         0.0         0.0           46.0         46.0         0.0         0.0         0.0         0.0         0.0           58.0         34.0         0.0         0.0         0.0         0.0         0.0           46.0         46.0         0.0         0.0         0.0         0.0         0.0           46.0         46.0         0.0         0.0         0.0         0.0         0.0           70.9         18.5         2.4         0.2         0.0         0.0         0.0           45.3         46.0         0.7         0.0         0.0         0.0         0.0           45.3         46.0         0.7         0.0         0.0         0.0         0.0           45.3         46.0         0.7         0.0         0.0         0.0         0.0           67.3         24.2         0.5         0.0         0.0         0.0         0.0           79.9         11.3         0.8         0.0         0.0         0.0         0.0           79.0         13.0         0.0	58         34         0         0         0         0         0           63.6         23.5         4.3         0.5         0.0         0.0         0.0           46.0         46.0         0.0         0.0         0.0         0.0         0.0           58.0         34.0         0.0         0.0         0.0         0.0         0.0           46.0         46.0         0.0         0.0         0.0         0.0         0.0           46.0         46.0         0.0         0.0         0.0         0.0         0.0           70.9         18.5         2.4         0.2         0.0         0.0         0.0           45.3         46.0         0.7         0.0         0.0         0.0         0.0           45.3         46.0         0.7         0.0         0.0         0.0         0.0           45.3         46.0         0.7         0.0         0.0         0.0         0.0           45.3         24.0         0.5         0.0         0.0         0.0         0.0           67.3         24.2         0.5         0.0         0.0         0.0         0.0           79.9 <td< td=""><td>58         34         0         0         0         0         0           63.6         23.5         4.3         0.5         0.0         0.0         0.0         10.1           46.0         46.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0           58.0         34.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         &lt;</td><td>58         34         0         0         0         0         0           63.6         23.5         4.3         0.5         0.0         0.0         0.0         10.1         1           46.0         46.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         1           58.0         34.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         1           46.0         46.0         0.0         0.0         0.0         0.0         0.0         0.0         6.3         1           0         1         2         3         4         5         5+         chi-sq         df           69         22         1         0         0         0         0         1.7         1           45.3         46.0         0.7         0.0         0.0         0.0         0.0         0.0         1           45.3         46.0         0.7         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.7         0           67.3         &lt;</td></td<>	58         34         0         0         0         0         0           63.6         23.5         4.3         0.5         0.0         0.0         0.0         10.1           46.0         46.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0           58.0         34.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         <	58         34         0         0         0         0         0           63.6         23.5         4.3         0.5         0.0         0.0         0.0         10.1         1           46.0         46.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         1           58.0         34.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         1           46.0         46.0         0.0         0.0         0.0         0.0         0.0         0.0         6.3         1           0         1         2         3         4         5         5+         chi-sq         df           69         22         1         0         0         0         0         1.7         1           45.3         46.0         0.7         0.0         0.0         0.0         0.0         0.0         1           45.3         46.0         0.7         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.0         0.7         0           67.3         <