

DORA-E-MON SGA

Arranged

1

Gtr I

T	1	0	2	0	2	0	1	0	2	0	1	0	2	0	1	0	2	0
A	0	0	2	0	2	0	0	0	2	0	0	0	2	0	0	0	2	0
B	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3

P P P P

5

T	3	1	0	2	0	1	3	1	0	2	0	1	3	1	0	2	0	1
A	0	0	0	2	0	1	0	0	0	2	0	1	0	0	0	2	0	1
B	3	3	2	3	3	3	3	3	2	3	3	3	3	H	3	3	3	3

P P P H

9

T	1	2	0	1	2	3	1	0	1	3	2	0	0	2	0	0	2	0
A	0	2	0	0	2	0	2	3	0	2	0	2	0	0	2	0	0	0
B	0	3	0	1	0	0	0	2	0	3	2	3	3	3	2	3	3	3

13

T	1	1	0	5	0	3	5	3	5	3	5	6	5	3	3	3	3	3
A	0	2	3	4	4	4	4	4	4	4	4	4	4	4	4	4	4	4
B	3	2	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3

17

T	5	5	5	5	8	5	1	0	2	0	1	3	3	3	0	0	0	0
A	7	5	5	5	5	5	0	2	0	1	0	0	0	0	0	0	0	0
B	0	5	5	5	5	5	0	2	0	1	0	0	0	0	0	0	0	0

P

21

T	1	1	0	5	0	3	5	3	3	5	3	5	6	5	3	3	3	3
A	0	2	3	4	4	4	4	4	4	4	4	4	4	4	4	4	4	4
B	3	2	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3	3

25

5 5 5 5 8 5 G 1 0 0 3 G 1 1 0 G 0 1 0

7 5 5 5 0 0 2 0 3 1 1 0 0 2 0

0 0 0 0 0 0 3 2 3 3 2 0 2 0

P

29

5 5 8 6 8 8 0 1 3 3 5 8 3 3 3 0 0 2 0 3 2 0 2

0 0 0 0 0 0 3 2 0 0 0 0 3 2 0 2

H

33

5 5 8 3 7 5 3 5 3 1 3 3 5 0 3 1 1 1 0 0

0 0 0 0 0 0 0 0 0 0 3 0 0 2 0 0 0

0 0 0 3 sl. sl. 3 sl. P

37

0 0 0 1 1 1 0 0 0 0 0 0 1 1 1 0 0

0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0

3 2 3 3 2 3 3 2 3 2 3 2

41

0 0 0 3 3 1 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

3 2 3 3 2 3 3 2 3 2 3 2