N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				
N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				
N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				
N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				
N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				
N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				
N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				
N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				
N:	5	10	15	20	25	30	35	40	45	50	55	60	65	70	75	80	85	90	95	100
Original	ATCCG	AATGG	CATGC	ATAGG	CCCAT	TAGCG	ATGCA	TGCCA	TTGCC	ATGCA	TTGAC	CATTG	CCGAT	TGAGC	ATTGC	TAAGC	ATTCG	CGATT	AGGCC	GATAG
Mutations																				