Theorem 5. The Hamming distance for χ^2 -statistics based on a 3×2 contingency table in GWAS is

$$= \frac{d\left(T_{3\times2}(p,q,r,s,t,u),T_{3\times2}(p',q',r',s',t',u')\right)}{\frac{|p-p'|+|q-q'|+|r-r'|+|s-s'|+|t-t'|+|u-u'|}{2}},$$

where $T_{3\times 2}(p,q,r,s,t,u)$ represents the following table data:

		Disease Status		Total
		0	1	IOtal
Genotype	0	p	q	p+q $r+s$
	1	r	s	
	2	t	u	t+u
Total		p+r+t	q+s+u	N

Proof. We consider the case where $(p>p') \wedge (q \leq q') \wedge (r \leq r') \wedge (s \leq s') \wedge (t \leq t') \wedge (u \leq u')$. Since p+q+r+s+t+u=p'+q'+r'+s'+t'+u'=N,

$$p - p' = (q' - q) + (r' - r) + (s' - s) + (t' - t) + (u' - u).$$

Thus, when we move p-p' elements out of p to q, r, \ldots, u by $(q'-q), (r'-r), \ldots, (u'-u)$, respectively, $T_{3\times 2}(p, q, r, s, t, u)$ changes to $T_{3\times 2}(p', q', r', s', t', u')$. Therefore, the Hamming distance dist satisfies

$$dist \leq p - p'$$

$$= \frac{|p - p'| + |q - q'| + |r - r'| + |s - s'| + |t - t'| + |u - u'|}{2}.$$

The similar discussions can be made for the other cases.

Here, when one element in $T_{3\times 2}(p,q,r,s,t,u)$ moves and the table changes to $T_{3\times 2}(\tilde{p},\tilde{q},\tilde{r},\tilde{s},\tilde{t},\tilde{u})$, the following inequality holds:

$$|p - \tilde{p}| + |q - \tilde{q}| + \dots + |u - \tilde{u}| \le 2.$$

Therefore, even when we move

$$k < \frac{|p - p'| + |q - q'| + |r - r'| + |s - s'| + |t - t'| + |u - u'|}{2}$$

elements and obtain $T_{3\times 2}(p'',q'',r'',s'',t'',u'')$, the table $T_{3\times 2}(p',q',r',s',t',u')$ never appears because

$$|p - p''| + |q - q''| + \dots + |u - u''|$$

 $\leq 2k < |p - p'| + |q - q'| + \dots + |u - u'|.$

Consequently, we can show that

$$dist = \frac{|p - p'| + |q - q'| + |r - r'| + |s - s'| + |t - t'| + |u - u'|}{2}.$$

Theorem 6. The Hamming distance for χ^2 -statistics based on a 2×2 contingency table in GWAS is

$$= \left\lceil \frac{d \left(T_{2 \times 2}(a,b,c,d), T_{2 \times 2}(a',b',c',d') \right)}{4} \right\rceil,$$

where $T_{2\times 2}(a,b,c,d)$ represents the following table data:

		Disease Status		Total
		0	1	Total
Allele	0	a	b	a+b
	1	c	d	c+d
Total		a+c	b+d	2N

Proof. We consider the case of $(a>a') \land (b \leq b') \land (c \leq c') \land (d \leq d')$. Since $a+b+c+d=a'+b'+c'+d', \ a-a'=(b'-b)+(c'-c)+(d'-d)$. Then, when we move a-a' elements out of a to b, c, and d by (b'-b), (c'-c), and (d'-d), respectively, $T_{2\times 2}(a,b,c,d)$ changes to $T_{2\times 2}(a',b',c',d')$. Therefore, the Hamming distance dist satisfies

$$\begin{array}{lcl} dist & \leq & \left\lceil \frac{a-a'}{2} \right\rceil \\ & = & \left\lceil \frac{|a-a'|+|b-b'|+|c-c'|+|d-d'|}{4} \right\rceil \end{array}$$

because a change in one individual of the dataset causes a change in two elements of the table. The similar discussions can be made for the other cases.

Here, in a similar manner to the proof of Theorem 5, we can show that when we move

$$k < \left\lceil \frac{|a - a'| + |b - b'| + |c - c'| + |d - d'|}{4} \right\rceil \tag{1}$$

elements from $T_{2\times 2}(a, b, c, d)$, the table $T_{2\times 2}(a', b', c', d')$ never appears, because $k \in \mathbb{N}_{>0}$ and (7) stands the following inequality:

$$k < \frac{|a-a'| + |b-b'| + |c-c'| + |d-d'|}{4}.$$

Consequently, the Hamming distance is

$$dist = \left\lceil \frac{|a - a'| + |b - b'| + |c - c'| + |d - d'|}{4} \right\rceil.$$