

# Supplementary Material for: Zero-Downtime Hardening of Legacy Patient Identifiers Against Transcription Errors: A Table-Free Verhoeff Formulation

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## ONLINE RESOURCE 1: WORKED EXAMPLE

Let the TAJ be  $\mathbf{t} = (0, 7, 6, 5, 4, 3, 2, 1, 0)$  in right-to-left order (i.e. the printed form is 012345670). We trace the TAJ-FC computation step by step.

$i$	$t_i$	$\pi_{i+1}(t_i)$	$S$ after $\text{Mul}(S, \varphi(\pi_{i+1}(t_i)))$	Note
0	0	1	(1, 0)	$\varphi(1) = (1, 0)$
1	7	5	(1, 1)	$\varphi(5) = (0, 1)$
2	6	5	(1, 0)	$\varphi(5) = (0, 1)$
3	5	3	(4, 0)	$\varphi(3) = (3, 0)$
4	4	7	(1, 1)	$\varphi(7) = (2, 1)$
5	3	5	(1, 0)	$\varphi(5) = (0, 1)$
6	2	9	(0, 1)	$\varphi(9) = (4, 1)$
7	1	5	(0, 0)	$\varphi(5) = (0, 1)$ ; back to identity
8	0	5	(0, 1)	$\varphi(5) = (0, 1)$ ; final state

Partial fold:  $F_{+1} = (0, 1)$ . Inverse:  $(0, 1)^{-1} = (0, 1)$  (a reflection is its own inverse). TAJ-FC:  $\varphi^{-1}(0, 1) = 0 + 5 = 5$ .

The receiver, given TAJ = 012345670 and TAJ-FC = 5, places the digit 5 at position 0 and folds the ten-digit sequence; the result is (0, 0), confirming validity.

Now suppose positions 4 and 5 (right-to-left) are transposed, changing the printed TAJ from 012345670 to 012354670. Recomputing the partial fold with the altered input yields  $F_{+1} = (0, 0)$  and therefore TAJ-FC = 0  $\neq$  5, triggering the hard stop.

## ONLINE RESOURCE 2: REFERENCE PSEUDOCODE

The two listings below translate the algebraic definitions into executable form. The group operations reduce to modulo-5 arithmetic plus a single XOR; the position-dependent permutation  $\pi_i$  uses two constant vectors  $\rho$ ,  $\rho^{-1}$  and modulo-10 arithmetic.

**Listing 1:** Primitive operations on  $S = \mathbb{Z}_5 \times \mathbb{Z}_2$  and the cycle-conjugate permutation  $\pi_i$ .

```
constant rho      := [0, 1, 4, 3, 8, 9, 6, 7, 2, 5]
constant rho_inv := [0, 1, 8, 3, 2, 9, 6, 7, 4, 5]

function Mul((a,b), (c,d)):
  sign := +1 if b = 0, else -1
  return ( (a + sign * c) mod 5,  b XOR d )

function Inv((a,b)):
  if b = 0: return ( (-a) mod 5, 0 )
  else:    return ( a, 1 )

function Phi(d):
  if d < 5: return (d, 0)
  else:    return (d - 5, 1)

function PhiInv((a,b)):
  if b = 0: return a
  else:    return a + 5
```

```

function Pi(d, i):                                -- position permutation
  x := rho_inv[d]
  y := (x + i) mod 10
  return rho[y]

```

**Listing 2:** TAJ-FC computation and vTAJ validation.

```

function ComputeTAJFC(t[0..8]):
  -- t[0] = rightmost digit
  S := (0, 0)                                     -- identity
  for i from 0 to 8:
    S := Mul(S, Phi(Pi(t[i], i + 1)))
  return PhiInv(Inv(S))

function ValidateVTAJ(t[0..8], k):
  -- k = received TAJ-FC
  S := (0, 0)
  S := Mul(S, Phi(Pi(k, 0)))                       -- position 0: TAJ-FC
  for i from 0 to 8:
    S := Mul(S, Phi(Pi(t[i], i + 1)))
  return S = (0, 0)

```

### ONLINE RESOURCE 3: ADDITIONAL MONTE CARLO DATA

Table S1 reports detection rates from the  $n = 8,000$  pilot run. Results are consistent with the  $n = 150,000$  run reported in the main text (Table I).

TABLE I  
\*

Table S1: Detection rates from Monte Carlo runs with  $n = 8,000$  trials per configuration (seed 1). Wilson 95% intervals in the last two

	Error model	Codeword family	$n$	Det.	Rate	$CI_{low}$	$CI_{high}$
	Single-digit subst.	legacy mod-10 (9)	8000	8000	1.0000	0.9995	1.0000
	Single-digit subst.	fold explicit (9)	8000	8000	1.0000	0.9995	1.0000
	Adjacent transpos.	legacy mod-10 (9)	8000	7059	0.8824	0.8751	0.8893
columns.	Adjacent transpos.	fold explicit (9)	8000	8000	1.0000	0.9995	1.0000
	Random noise	legacy mod-10 (9)	8000	7543	0.9429	0.9376	0.9478
	Random noise	fold explicit (9)	8000	8000	1.0000	0.9995	1.0000
	Single-digit subst.	full vTAJ (10)	8000	8000	1.0000	0.9995	1.0000
	Adjacent transpos.	full vTAJ (10)	8000	8000	1.0000	0.9995	1.0000
	Random noise	full vTAJ (10)	8000	8000	1.0000	0.9995	1.0000

Figure S1 shows the convergence of Monte Carlo estimates between the two sample sizes. Point estimates are stable; Wilson intervals narrow by a factor of  $\sqrt{150,000/8,000} \approx 4.3$ , as expected from standard sampling theory.

Monte Carlo stability: same configuration at two sample sizes (95% Wilson intervals)

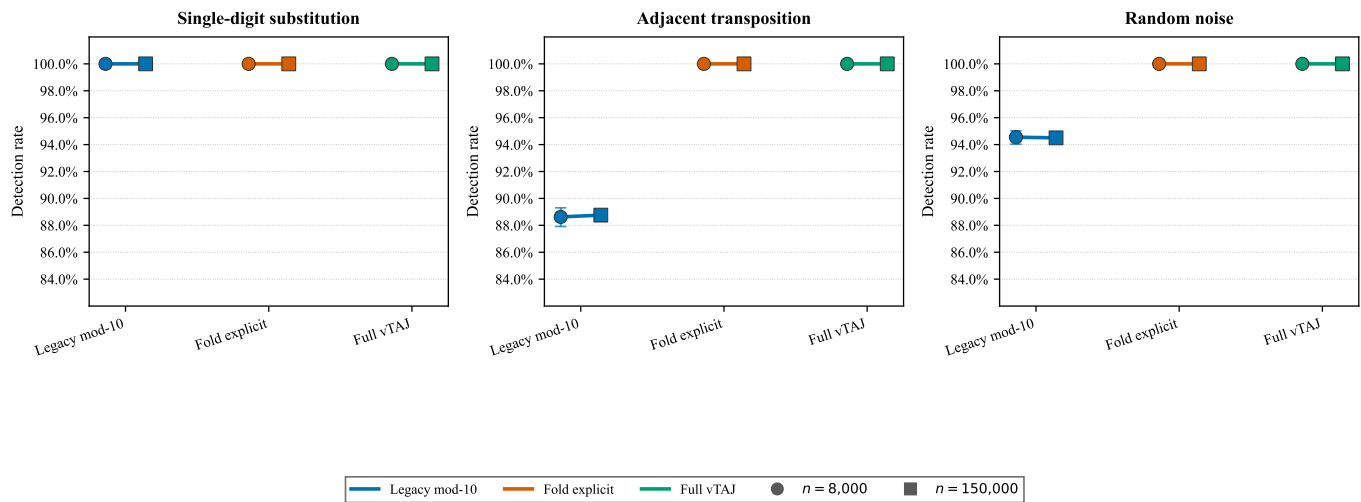


Fig. 1. \*

Figure S1: Convergence of Monte Carlo estimates between  $n = 8,000$  (circles) and  $n = 150,000$  (squares). Thin horizontal ticks show Wilson 95% interval endpoints.