

**Supplementary Table 1. Estimated ERSPC model parameters\***

<b>Onset is modeled by a hazard function (5 parameters)†</b>			<b>Transition probability (12 parameters)</b>		
Age in years	Hazard		From	To	Probability
0	0.0000		T1G6Mx	T1G7Mx	0.27
30	0.0000			T2G6Mx	0.71
50	0.0005			Clinical	0.02
70	0.0170		T1G7Mx	T1G8Mx	0.19
100	0.0503			T2G7Mx	0.78
				Clinical	0.03
			T1G8Mx	T2G8Mx	0.00
				Clinical	1.00
<b>Durations in preclinical states are modeled by Weibull distributions (13 parameters)</b>			T2G6Mx	T2G7Mx	0.24
	Mean	Shape		T3G6Mx	0.62
T1G6M0	2.70868	0.90514		Clinical	0.15
T1G7M0	6.49444	0.90514	T2G7Mx	T2G8Mx	0.63
T1G8M0	21.6563	0.90514		T3G7Mx	0.00
T2G6M0	4.00188	3.84944		Clinical	0.36
T2G7M0	7.40624	3.84944	T2G8Mx	T3G8Mx	0.00
T2G8M0	11.1226	3.84944		Clinical	1.00
T3G6M0	3.53881	1.04132	T3G6Mx	T3G7Mx	0.00
T3G7M0	18.9094	1.04132		Clinical	1.00
T3G8M0	23.3957	1.04132	T3G7Mx	T3G8Mx	0.00
TxGxM1 = mean(TxGxM1) × 0.0101638				Clinical	1.00
			T3G8Mx	Clinical	1.00
<b>Correlation between the duration of preclinical states in an individual (1 parameter)</b>			<b>Yearly hazard of additional clinical diagnosis in different preclinical states (4 parameters)‡</b>		
Correlation 0.866907			T1GxM0		0.056
<b>Yearly hazard of metastasis in different preclinical states (9 parameters)</b>			T2GxM0		0.001
T1G6M0	0.0003		T3GxM0		0.056
T1G7M0	0.0016		T1GxM1		0.801
T1G8M0	0.0213		T2GxM1		0.014
T2G6M0	0.0045		T3GxM1		0.810
T2G7M0	0.0119		<b>Sensitivity (4 parameters)</b>		
T2G8M0	0.0473		T1GxM0		0.936
T3G6M0	0.0017		T2GxM0		0.941
T3G7M0	0.0067		T3GxM0		1.000
T3G8M0	0.1046		T1GxM1		0.962
			T2GxM1		0.965
			T3GxM1		1.000

\*T1, T2, T3 are the three clinical T stages: T1 (impalpable), T2 (palpable, confined to the prostate), and T3+ (palpable, with extensions beyond the prostatic capsule), G6 is Gleason score (G) less than 7, G7 is Gleason score of 7, G8 Gleason score more than 7, M0 is the local-regional stage and M1 is the distant stage. Tx, Gx, and Mx imply all clinical T stages, all Gleason scores, and all metastatic stages, respectively.

†Time of tumor onset is generated from a piece-wise linear hazard function.

‡These parameters model a higher incidence in the control arm of the ERSPC trial compared to the population in 1991 (the baseline). The difference in clinical diagnosis between these two populations can for instance be attributed to contamination (screening in the control arm) or to changes in clinical practice leading to earlier diagnosis, eg, to the use of PSA testing for symptomatic disease in a clinical setting.

**Supplementary Table 2.** Observed data in ERSPC–Rotterdam\*

<b>Baseline</b>							
Age group, y		n	Incidence per 1000 man-years †				
45–49		11	0.02				
50–54		58	0.14				
55–59		131	0.36				
60–64		387	1.19				
65–69		734	2.59				
70–74		950	4.50				
75–79		969	6.57				
80–84		668	7.98				
>85		431	8.52				
Clinical T stage and metastatic stage distribution, % (n = 1610)‡							
		T1	T2	T3	Total		
M0		12.92	47.45	17.76	78.14		
M1		1.06	11.30	9.50	21.86		
Total		13.98	58.76	27.27	100.00		
<b>Control arm§</b>							
Age group, y		n	Incidence per 1000 man-years				
55–59		29	1.61				
60–64		135	3.35				
65–69		192	5.13				
70–74		247	7.95				
>75		146	9.85				
Clinical T stage, metastatic stage, and Gleason score distribution, % (n = 472)							
				G < 7	G = 7	G > 7	Total
M0	T1			29.87	7.84	4.45	42.16
	T2			15.47	10.59	3.81	29.87
	T3			4.24	9.32	7.20	20.76
	Subtotal			49.58	27.75	15.47	92.80
M1	T1			0.00	0.00	0.00	0.00
	T2			0.21	0.85	1.69	2.75
	T3			0.00	1.27	3.18	4.45
	Subtotal			0.21	2.12	4.87	7.20
Total				49.79	29.87	20.34	100.00
<b>Screen arm§</b>							
Detection rate per 1000 men screened by round of screening							
		Round 1		Round 2		Round 3	
Age group, y	n	rate	n	rate	n	rate	
55–59	176	27.64	20	20.77	0	0.00	
60–64	239	44.81	171	35.14	38	26.44	
65–69	366	76.81	189	49.74	83	32.35	
70–74	287	86.73	170	59.30	74	37.85	
Clinical T stage, metastatic stage, and Gleason score distribution, % (n = 1815)							
Round 1				G < 7	G = 7	G > 7	Total
M0	T1			28.23	6.61	1.04	35.88
	T2			29.84	12.75	3.59	46.18
	T3			6.89	7.84	2.64	17.37
	Subtotal			64.97	27.20	7.27	99.43
M1	T1			0.00	0.00	0.00	0.00
	T2			0.00	0.00	0.00	0.00
	T3			0.00	0.09	0.47	0.57

	Subtotal	0.00	0.09	0.47	0.57
	Total	64.97	27.29	7.74	100.00
Round 2		G < 7	G = 7	G > 7	Total
M0	T1	52.00	9.09	0.73	61.82
	T2	24.73	7.64	1.82	34.18
	T3	2.00	1.27	0.36	3.64
	Subtotal	78.73	18.00	2.91	99.64
M1	T1	0.00	0.00	0.00	0.00
	T2	0.18	0.00	0.18	0.36
	T3	0.00	0.00	0.00	0.00
	Subtotal	0.18	0.00	0.18	0.36
Total		78.91	18.00	3.09	100.00
Round 3		G < 7	G = 7	G > 7	Total
M0	T1	56.41	7.18	0.00	63.59
	T2	27.69	4.62	2.56	34.87
	T3	0.51	0.00	1.03	1.54
	Subtotal	84.62	11.79	3.59	100.00
M1	T1	0.00	0.00	0.00	0.00
	T2	0.00	0.00	0.00	0.00
	T3	0.00	0.00	0.00	0.00
	Subtotal	0.00	0.00	0.00	0.00
Total		84.62	11.79	3.59	100.00

Incidence of interval cancers per 1000 man-years by time since last screening

Time (t) since last screening, y	n	Incidence
0 ≤ t < 1	8	0.50
1 ≤ t < 2	7	0.48
2 ≤ t < 3	12	0.84
3 ≤ t < 4	18	1.46
t ≥ 4	52	4.21

\*T1, T2, T3 are the three clinical T stages: T1 (impalpable), T2 (palpable, confined to the prostate), and T3+ (palpable, with extensions beyond the prostatic capsule), G < 7 is Gleason score (G) less than 7, G = 7 is Gleason score of 7, G > 7 Gleason score more than 7, M0 is the local or regional stage and M1 is the distant stage. For clinical T stage, metastatic stage, and Gleason score distribution, only completed cases (cases for which the clinical T stage, Gleason score and metastatic stage are known) are shown.

†Data from the Netherlands Cancer Registry for 1991.

‡Data from the Rotterdam Cancer Registry for 1992 and 1993.

§ For the screen arm, data for cases diagnosed up to December 31, 2006, are presented. For the control arm, data for cases diagnosed up to July 4, 2004, are presented. Data for the control arm were collected from the Rotterdam cancer registry. Because there is a delay of 1–2 years in the reporting of tumors, the cutoff date for the control arm is different from that of the screen arm.

**Supplementary Table 3. Observed incidence and stage distribution in the US population\***

Year	Age group (years)						
	50–54	55–59	60–64	65–69	70–74	75–79	80–84
<b>Observed incidence per 1000 man-years</b>							
1985	0.27	0.94	2.26	4.48	6.81	9.15	11.09
1986	0.32	0.93	2.34	4.79	7.09	9.47	11.60
1987	0.37	1.16	2.61	5.29	8.25	10.90	12.14
1988	0.37	1.11	2.81	5.43	8.86	10.91	12.81
1989	0.36	1.21	3.16	6.15	9.19	11.58	12.82
1990	0.49	1.42	3.66	7.24	10.76	13.39	14.79
1991	0.70	1.93	4.93	9.36	14.47	16.78	17.47
1992	0.90	2.57	6.04	11.31	15.80	18.14	17.75
1993	0.95	2.71	5.97	10.66	13.92	14.45	13.99
1994	0.99	2.60	5.54	9.33	12.16	11.76	10.65
1995	1.04	2.85	5.39	8.71	11.14	10.55	9.65
1996	1.19	2.85	5.78	8.34	10.65	10.44	9.00
1997	1.27	3.04	5.72	9.00	10.80	10.47	9.38
1998	1.27	3.15	5.49	8.69	10.36	10.79	9.16
1999	1.40	3.29	6.28	9.41	11.37	10.96	9.12
2000	1.39	3.55	6.17	9.35	10.75	10.70	9.25
<b>Observed percentage of distant disease</b>							
1985	18.85	18.84	17.51	19.00	19.41	21.04	24.34
1986	18.75	19.86	16.78	15.77	19.89	20.04	23.93
1987	21.30	15.71	14.31	14.86	15.50	17.48	22.33
1988	23.84	15.29	14.83	14.87	15.20	17.56	21.11
1989	15.88	15.41	13.79	14.12	13.51	17.41	22.55
1990	11.59	13.39	12.10	11.44	12.28	14.94	19.45
1991	7.62	8.01	8.45	8.36	9.44	11.71	19.17
1992	5.69	6.47	6.23	5.78	7.12	9.53	12.99
1993	4.52	5.36	5.98	5.30	6.04	8.92	14.49
1994	2.27	4.81	4.53	5.38	5.27	8.26	13.70
1995	4.01	3.62	5.13	4.66	6.22	8.31	12.95
1996	4.18	3.82	3.69	3.71	5.29	7.49	13.18
1997	3.65	2.91	3.35	4.13	4.87	6.39	11.52
1998	2.33	3.30	3.70	3.31	4.75	6.62	11.31
1999	3.16	2.62	2.67	3.65	3.99	5.21	9.59
2000	2.00	2.58	2.14	3.23	3.84	5.60	8.89

\*Data from the National Cancer Institute's Surveillance, Epidemiology, and End Results database. The data are based on nine catchment areas of the United States: San Francisco–Oakland, Connecticut, Detroit, Hawaii, Iowa, New Mexico, Seattle, Utah, and Atlanta.