

**Supplemental Table 1. Semen parameters and cancer risk in men with SA versus fertile men**

	<u>All-Site</u>		<u>Prostate</u>		<u>Melanoma</u>		<u>Testicular</u>		<u>Other</u>	
	HR	95% CI	HR	95% CI	HR	95% CI	HR	95% CI	HR	95% CI
<b>Model 1: Infertile vs. Fertile Controls</b>										
Infertile	1.17	(0.96, 1.41)	0.90	(0.54, 1.52)	1.35	(0.9, 2.02)	3.30	(1.59, 6.86)	0.99	(0.76, 1.29)
<b>Model 2: Concentration vs. Fertile Controls</b>										
Azoospermia	1.00	(0.49, 2.06)	0.37	(0.04, 3.26)	0.90	(0.19, 4.25)	3.67	(0.45, 30.25)	1.12	(0.44, 2.85)
Oligozoospermic	1.66	(1.13, 2.41)	0.53	(0.12, 2.26)	1.12	(0.44, 2.88)	11.89	(4.91, 28.82)	1.36	(0.79, 2.35)
Normozoospermic	1.08	(0.84, 1.39)	1.12	(0.59, 2.12)	0.96	(0.55, 1.68)	2.86	(1.22, 6.71)	0.97	(0.69, 1.36)
Hyperzoospermic	1.15	(0.88, 1.50)	0.87	(0.39, 1.97)	1.95	(1.21, 3.16)	1.60	(0.57, 4.45)	0.90	(0.62, 1.31)
<b>Model 3: Count vs. Fertile Controls</b>										
Azoospermia	1.01	(0.49, 2.08)	0.38	(0.04, 3.32)	0.92	(0.19, 4.31)	3.51	(0.42, 29.3)	1.15	(0.45, 2.92)
Oligozoospermic	1.78	(1.23, 2.58)	0.88	(0.28, 2.77)	1.29	(0.54, 3.11)	10.33	(4.07, 26.23)	1.59	(0.95, 2.65)
Normozoospermic	1.09	(0.79, 1.5)	1.12	(0.52, 2.39)	1.01	(0.48, 2.1)	3.53	(1.32, 9.41)	0.92	(0.58, 1.44)
Hyperzoospermic	1.10	(0.88, 1.50)	0.87	(0.43, 1.74)	1.52	(0.97, 2.38)	2.17	(0.94, 5.01)	0.91	(0.66, 1.25)
<b>Model 4: Motility vs. Fertile Controls</b>										
Azoospermia	1.00	(0.48, 2.05)	0.38	(0.04, 3.35)	0.92	(0.2, 4.32)	3.60	(0.43, 29.81)	1.12	(0.44, 2.85)
Q1 (0-49)	1.50	(1.13, 1.99)	1.37	(0.69, 2.71)	1.60	(0.89, 2.89)	4.12	(1.53, 11.09)	1.31	(0.88, 1.94)
Q2 (50-59)	1.31	(0.98, 1.75)	1.18	(0.51, 2.73)	1.24	(0.66, 2.35)	4.23	(1.68, 10.64)	1.14	(0.76, 1.7)
Q3 (60-69)	0.86	(0.61, 1.22)	0.26	(0.06, 1.1)	1.22	(0.63, 2.34)	2.19	(0.74, 6.53)	0.79	(0.49, 1.27)
Q4 (70-100)	0.96	(0.66, 1.4)	0.80	(0.17, 3.86)	1.43	(0.72, 2.85)	2.87	(1.06, 7.79)	0.64	(0.36, 1.14)
<b>Model 5: Vitality vs. Fertile Controls (UU Sample Only)</b>										
Azoospermia	1.28	(0.6, 2.74)	0.50	(0.05, 4.74)	1.55	(0.34, 7.1)	3.92	(0.45, 33.88)	1.28	(0.46, 3.56)
Q1 (0-45)	1.40	(1.02, 1.92)	1.25	(0.6, 2.57)	1.26	(0.63, 2.51)	6.63	(2.45, 17.94)	1.13	(0.71, 1.82)
Q2 (46-55)	1.16	(0.82, 1.64)	1.58	(0.7, 3.6)	0.98	(0.45, 2.14)	1.15	(0.24, 5.56)	1.15	(0.72, 1.83)
Q3 (56-64.4)	0.94	(0.65, 1.36)	0.18	(0.02, 1.36)	0.72	(0.3, 1.72)	2.75	(0.93, 8.19)	1.04	(0.65, 1.65)
Q4 (64.5-94)	0.79	(0.51, 1.22)	*		1.87	(0.97, 3.61)	0.61	(0.07, 4.93)	0.55	(0.28, 1.06)

**Model 6: Head Morphology vs. Fertile Controls (UU Sample Only)**

Azoospermia	1.28	(0.6, 2.73)	0.52	(0.05, 4.95)	1.51	(0.33, 6.96)	3.92	(0.44, 34.99)	1.26	(0.45, 3.51)
Q1	1.09	(0.75, 1.58)	1.02	(0.4, 2.58)	0.63	(0.25, 1.63)	4.22	(1.43, 12.5)	1.07	(0.65, 1.77)
Q2	1.06	(0.75, 1.51)	1.28	(0.56, 2.95)	1.07	(0.51, 2.24)	0.99	(0.21, 4.77)	1.03	(0.64, 1.64)
Q3	1.06	(0.74, 1.51)	0.95	(0.38, 2.39)	1.13	(0.54, 2.38)	1.96	(0.51, 7.55)	1.00	(0.62, 1.62)
Q4	0.96	(0.65, 1.41)	0.34	(0.08, 1.48)	1.83	(0.97, 3.44)	2.10	(0.62, 7.14)	0.68	(0.38, 1.23)

**Model 7: Tail Morphology vs. Fertile Controls (UU Sample Only)**

Azoospermia	1.28	(0.6, 2.74)	0.52	(0.05, 5.01)	1.53	(0.33, 6.99)	4.15	(0.46, 37.03)	1.26	(0.45, 3.52)
Q1	1.21	(0.86, 1.71)	1.42	(0.67, 2.99)	1.18	(0.56, 2.5)	2.86	(0.83, 9.87)	1.03	(0.62, 1.7)
Q2	1.27	(0.91, 1.77)	1.26	(0.53, 2.99)	1.16	(0.57, 2.36)	5.31	(1.91, 14.77)	1.04	(0.64, 1.67)
Q3	1.01	(0.7, 1.46)	0.44	(0.1, 1.95)	1.61	(0.86, 3.04)	0.48	(0.06, 3.9)	0.96	(0.58, 1.57)
Q4	0.66	(0.42, 1.03)	0.31	(0.07, 1.35)	0.57	(0.2, 1.62)	1.12	(0.23, 5.4)	0.77	(0.44, 1.33)

**Model 8: Total Motile Count vs. Fertile Controls (UU Sample Only)**

Azoospermia	1.29	(0.6, 2.75)	0.52	(0.05, 4.99)	1.56	(0.34, 7.17)	3.65	(0.42, 31.76)	1.29	(0.46, 3.59)
Q1	1.58	(1.17, 2.13)	1.16	(0.54, 2.5)	0.99	(0.47, 2.09)	6.85	(2.67, 17.59)	1.56	(1.03, 2.37)
Q2	0.69	(0.46, 1.04)	0.90	(0.38, 2.15)	0.59	(0.23, 1.51)	1.83	(0.47, 7.07)	0.58	(0.32, 1.05)
Q3	1.04	(0.72, 1.51)	0.86	(0.25, 2.93)	1.41	(0.72, 2.75)	1.60	(0.42, 6.16)	0.88	(0.52, 1.5)
Q4	1.02	(0.69, 1.51)	0.45	(0.1, 2.01)	1.97	(1, 3.87)	0.55	(0.07, 4.48)	0.90	(0.53, 1.53)