## Test for shape dependence for the HSCT data

Covariates available in the dataset are:

 $\mathbf{age}$  scaled

race 1 = white, 0 = otherwise

 $\mathbf{gender}\ 1 = \mathrm{male}$ 

**allo** 1 = allogeneic, 0 = autologous

 $\mathbf{lym}\ 1 = \mathrm{lymphomas}\ \mathrm{disease}\ \mathrm{at}\ \mathrm{transplant}$ 

**heme1** 1 = heme at remission, 0 = otherwise

**heme2** 1 = heme at relapse, 0 = otherwise

cmv1 1 = receiver negative, 0 = otherwise

 $\mathbf{cmv2} \ 1 = \mathbf{receiver} \ \mathbf{and} \ \mathbf{donor} \ \mathbf{both} \ \mathbf{negative}, \ 0 = \mathbf{otherwise}$ 

Table 1: Different combinations and the testing results. The p-value is approximated with n = 100 bootstrap samples.

	covariates									
age	race	gender	allo	lym	heme1	heme2	cmv1	cmv2	p	
<b>√</b>			✓						0.63	
	✓		✓						0.11	
		✓	✓						0.14	
			✓	✓					0.41	
			✓		✓				0.35	
			✓			✓			0.42	
			✓				✓		0.32	
			✓					✓	0.08	

Table 2: Different combinations and the testing results. The p-value is approximated with n=100 bootstrap samples.

	covariates								
age	race	gender	allo	lym	heme1	heme2	cmv1	cmv2	p
$\checkmark$	✓		✓						0.46
$\checkmark$		✓	✓						0.34
	✓	✓	✓						0.05
		✓	✓	✓					0.06
$\checkmark$			✓	✓					0.67
		✓	✓					✓	0.10
		✓	✓			✓			0.17
$\overline{\hspace{1em}}$			✓		✓				0.52
$\checkmark$			✓			✓			0.46
$\checkmark$			✓				✓		0.74
$\checkmark$			✓					✓	0.22
$\overline{\hspace{1em}}$		✓	✓						0.32
		✓	✓		✓				0.34
		✓	✓			✓			0.35
		✓	✓				✓		0.28
		✓	✓					✓	0.09
			✓		✓			✓	0.09
			✓		✓		✓		0.58
			✓			✓		✓	0.20
			✓			✓	✓		0.54

Table 3: Different combinations and the testing results. The p-value is approximated with n = 100 bootstrap samples.

				iates	covar				
	cmv2	cmv1	heme2	heme1	lym	allo	gender	race	age
0.					✓	✓	✓	✓	
0.					✓	✓	✓		✓
0.						✓	✓	✓	✓
0.	✓		✓			✓	✓		
0.					✓	✓		✓	✓
0.	✓		✓			✓		✓	
0.	✓		✓			✓			✓
0.	✓					✓	✓		<b>√</b>
0.	✓		✓		✓	✓			
0.			✓			✓	✓	✓	
0.	✓					✓	✓	✓	
0.		✓		✓		✓			<b>√</b>
0.				✓	✓	✓	✓		
0.				✓		✓	✓		<b>√</b>
0.				✓		✓	✓	✓	
0.	✓		✓			✓	✓		<b>√</b>
0.	✓				✓	✓	✓		
0.	✓			✓		✓	✓		
0.	✓				✓	✓	✓		
0.	✓					✓	✓		<b>√</b>
0.	<b>√</b>					<b>√</b>	<b>√</b>	<b>√</b>	

Table 4: Different combinations and the testing results. The p-value is approximated with n = 100 bootstrap samples.

	covariates										
age	race	gender	allo	lym	heme1	heme2	cmv1	cmv2	p		
<b>√</b>		✓	✓		✓			✓	0.48		
		✓	✓	✓	✓			✓	0.28		
	✓	✓	✓			✓		✓	0.38		
<b>√</b>		✓	✓			✓		✓	0.37		