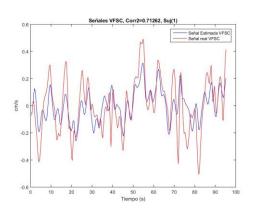
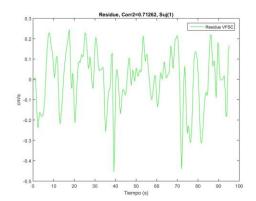
1 Residuos Modelos ARX pacientes TEC (PAM, PIC, PCC – VFSC)

A continuación se presentan los residuos de los sujetos con TEC modelados con SVM mediante modelos ARX. El primer gráfico representa la señal real de la velocidad del flujo sanguíneo del paciente y la estimada por el modelo, de color rojo y azul respectivamente. El segundo gráfico representa el residuo que existe entre ambas señales

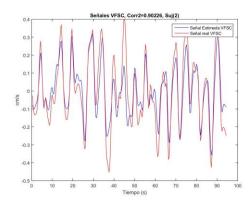
1.1 Sujeto 1

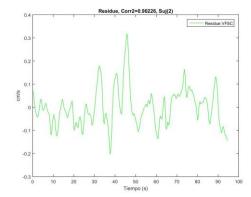
Correlación: 0.713





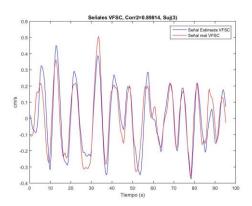
1.2 Sujeto 2

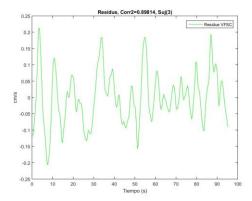




1.3 Sujeto 3

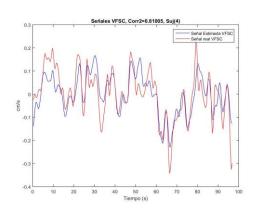
Correlación: 0.898

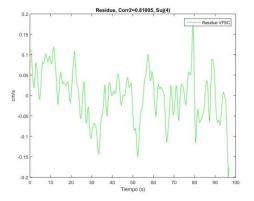




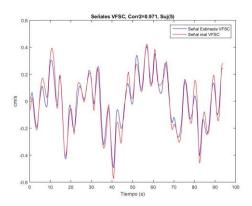
1.4 Sujeto 4

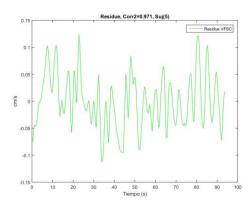
Correlación: 0.81





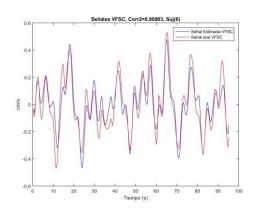
1.5 Sujeto 5

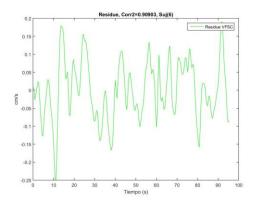




1.6 Sujeto 6

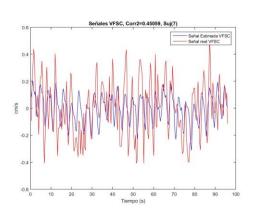
Correlación: 0.909

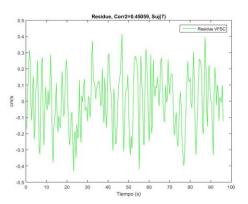




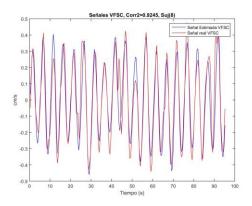
1.7 Sujeto 7

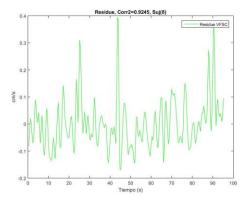
Correlación: 0.456





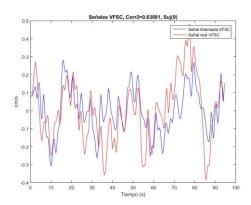
1.8 Sujeto 8

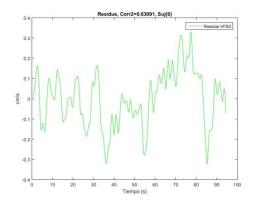




1.9 Sujeto 9

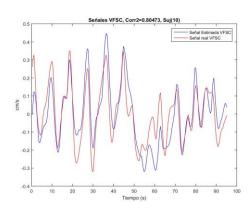
Correlación: 0.631

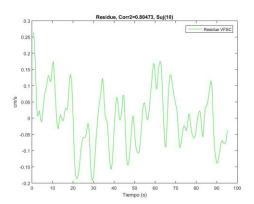




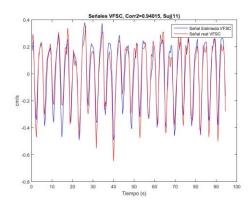
1.10 Sujeto 10

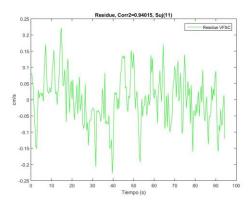
Correlación: 0.805





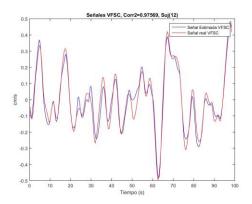
1.11 Sujeto 11

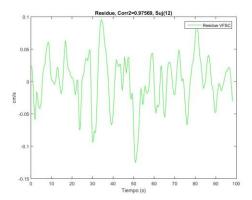




1.12 Sujeto 12

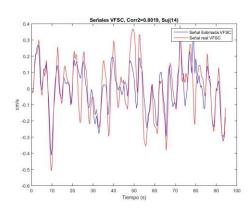
Correlación: 0.976

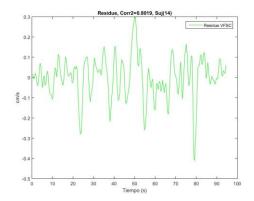




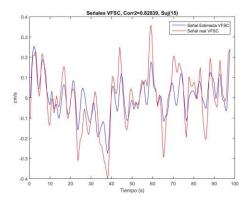
1.13 Sujeto 13

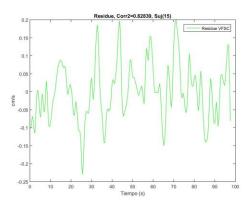
Correlación: 0.802





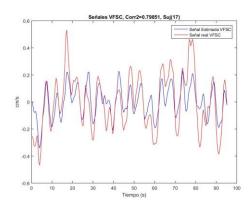
1.14 Sujeto 14

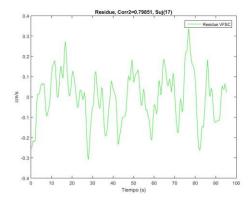




1.15 Sujeto 15

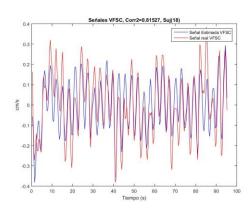
Correlación: 0.799

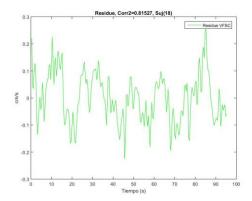




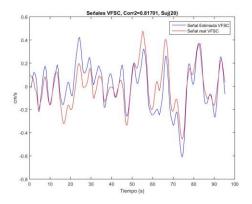
1.16 Sujeto 16

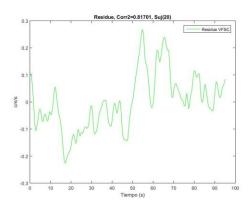
Correlación: 0.815





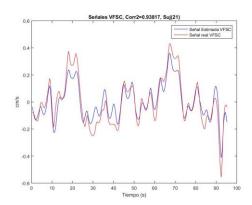
1.17 Sujeto 17

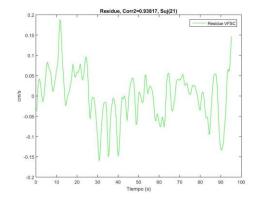




1.18 Sujeto 18

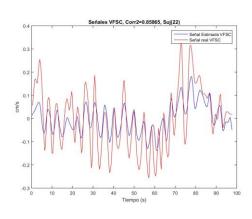
Correlación: 0.938

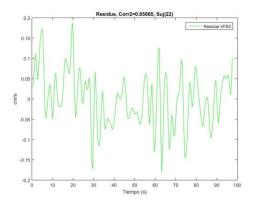




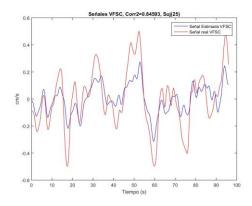
1.19 Sujeto 19

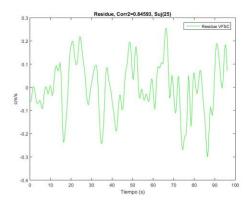
Correlación: 0.859





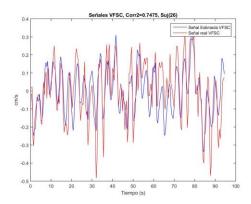
1.20 Sujeto 20

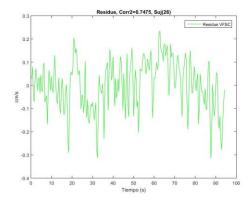




1.21 Sujeto 21

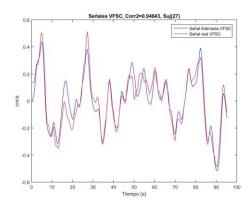
Correlación: 0.748

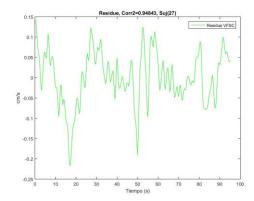




1.22 Sujeto 22

Correlación: 0.948





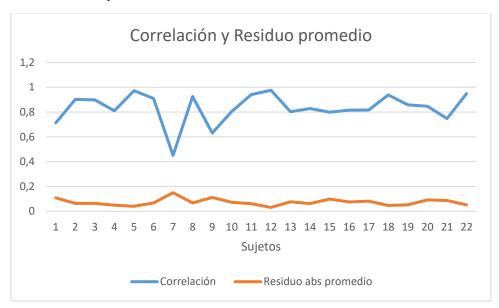
2 Promedio señales

A continuación se verá el comportamiento de las señales a modo general de la correlación de cada modelo en comparación con su residuo de acuerdo al modelo generado.

Sujeto Con	rrelación	Residuo abs promedio
1 0.7	126	0.1086
2 0.90	023	0.063
3 0.89	981	0.0634
4 0.8	101	0.0492
5 0.9	710	0.0396
6 0.90	090	0.0672
7 0.43	506	0.1496
8 0.92	245	0.0666
9 0.63	309	0.1108

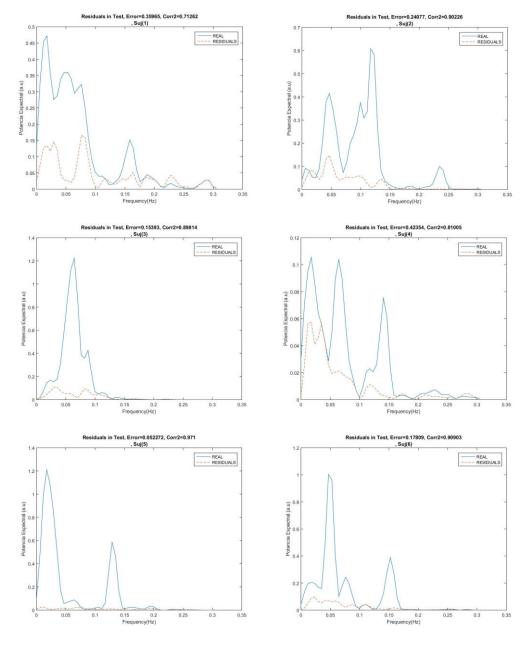
10	0.8047	0.0729
11	0.9402	0.0607
12	0.9757	0.0309
13	0.8019	0.0762
14	0.8284	0.0616
15	0.7985	0.0984
16	0.8153	0.0748
17	0.8170	0.0809
18	0.9382	0.0467
19	0.8586	0.0523
20	0.8459	0.0913
21	0.7475	0.0871
22	0.9484	0.0515
Promedio	0.833	0.073

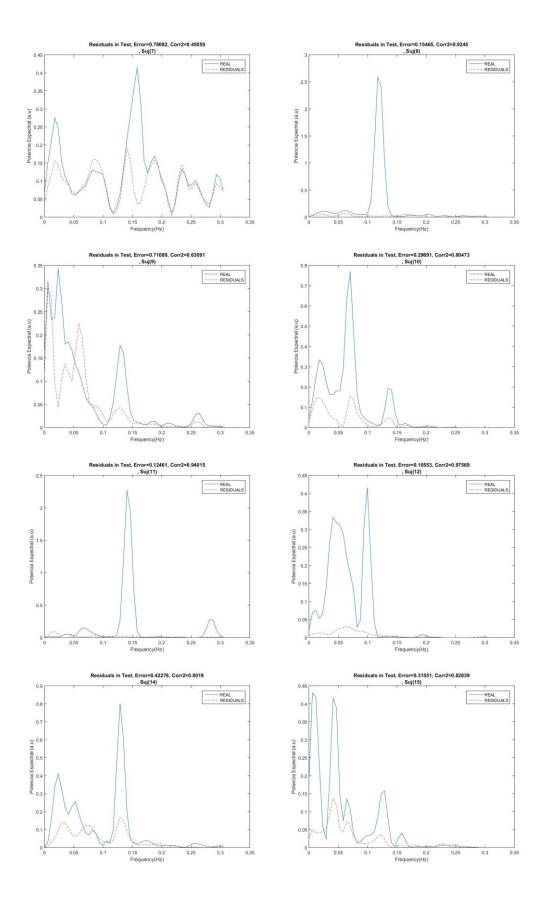
Se puede observar que el sujeto que presenta una menor correlación es el número 7 (GREG2000) con un valor cercano a 0.45 y un residuo promedio superior a 1, y el que presenta una mayor correlación es el número 12 () con un valor cercano a 0.97 y un residuo de 0.03.

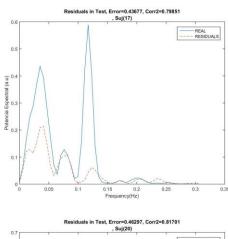


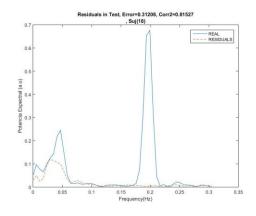
3 Potencia espectral de los modelos

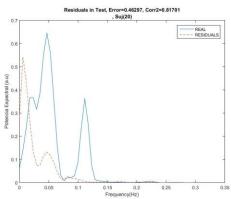
A continuación se encuentran los gráficos que representan la potencia espectral de la señal real y residual de los sujetos, en orden.

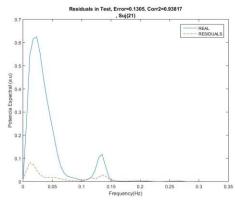


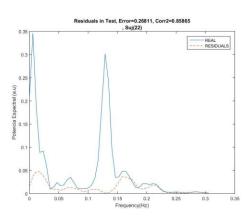


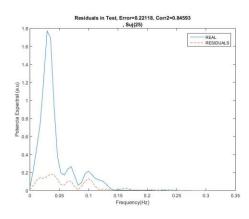


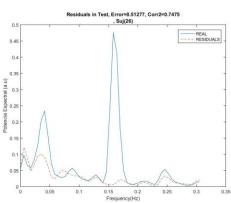


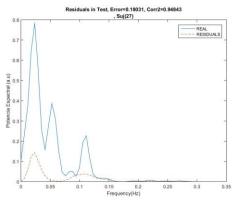












3.1 Promedio de la potencia espectral de los sujetos

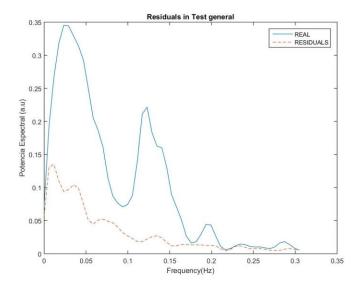


Ilustración 1 Residuos promedio del modelo multivariado no lineal ARX PAM, PIC, PCC - VFSC