

Imagerie Cellulaire et High Content Screening 11 Octobre 2018 – de 9h00 à 17h00

PROGRAMME				
09:00	Café d'accueil			
09:15	Introduction			
09:30	PRESENTATION	Hands-on 3D samples Analysis with Harmony 4.8	Alexander Schreiner	PerkinElmer
10:30	PAUSE			
11:00	PRESENTATION	Analysis of mitochondrial network morphology in Parkinson's disease cell models.	Nathalie Geyer	Sanofi, Strasbourg
11:40	PRESENTATION	Big Data Management from Organ to Cell level in HCA projects	Anne Danckaert	Institut Pasteur, Paris
12:20	Déjeuner			
13:40	PRESENTATION	Improving hit selection criteria during drug discovery process through High Content Screening	Benjamin Chanrion	Institut de Recherches Servier
14:20	PRESENTATION	Developing high-content imaging quantification of mitochondrial network morphology in fibroblasts from patients with inherited mitochondrial diseases	Aurélien Bayot	Institut Pasteur, Paris
15:00	PAUSE			
15:30	PRESENTATION	Use of Cell Painting to assess quality and diversity of an annotated chemical library	Julien Bellis	Evotec
16:10	PRESENTATION	Signals Screening – PerkinElmer's intuitive, app- based analysis tool for HTS and HCS data analysis	Christian Schueller	PerkinElmer
17:00	CONCLUSION			

Nous vous souhaitons un bon séminaire

