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Nota di bibliografia	Includes bibliographical references.
Nota di contenuto	Part 1 - Computational Methods A Review of Computational Methods for Molecular Imaging, by Fei Gao, Pengcheng Shi Fuzzy Connectedness Image Co-Segmentation for Hybrid PET/MRI and PET/CT Scans, by Ziyue Xu, Ulas Bagci, Jayaram Udupa, Daniel Mollura PET/MRI/VCT: Restoration of Virtual CT from Transmission Scan on PET/MRI using Joint-Anisotropic Diffusion, by Kuangyu Shi, Xiaoyin Cheng, Nassir Navab, Stefan Foerster, Sibylle I. Ziegler Large Scale Simplex Optimisation to Accelerate Kinetic Analysis, by Nicholas Dowson, Paul Thomas, Jye Smith, Olivier Salvado, Stephen Rose Gradient Projection for Regularized Cryo-Electron Tomographic Reconstruction, by Shadi Albargouni, Tobias Lasser, Weaam Alkhaldi,

	Ashraf Al-Amoudi, Nassir Navab Joint direct motion
	estimation/kinetic images reconstruction from gated PET data, by
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	Soft-Tissue from Single Short-TE MR Imaging Modality, by Anahita
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	by Xiaoyan Shen, Zhiliang Liu, Zhenghui Hu, Huafeng Liu
	Investigation of Single- Versus Joint-Modality PET-MR Registration for
	18F-Florbetapir Quantification: Application to Alzheimer's Disease, by
	Liam Cattell, Julia Schnabel, Jerome Declerck, Chloe Hutton.
Sommario/riassunto	This volume contains original submissions on the development and
	application of molecular imaging computing. The editors invited
	authors to submit high-quality contributions on a wide range of topics
	including, but not limited to: • Image Synthesis & Reconstruction of
	Emission Tomography (PET, SPECT) and other Molecular Imaging
	Modalities • Molecular Imaging Enhancement • Data Analysis of Clinical
	& Pre-clinical Molecular Imaging • Multi-Modal Image Processing
	(PET/CT. PET/MR. SPECT/CT. etc.) • Machine Learning and Data Mining
	in Molecular Imaging. Molecular imaging is an evolving clinical and
	research discipline enabling the visualization, characterization and
	guantification of biological processes taking place at the cellular and
	subcellular levels within intact living subjects. Computational methods
	play an important role in the development of molecular imaging, from
	image synthesis to data analysis and from clinical diagnosis to therapy
	individualization. This work will bring readers from academia and
	industry up to date on the most recent developments in this field.