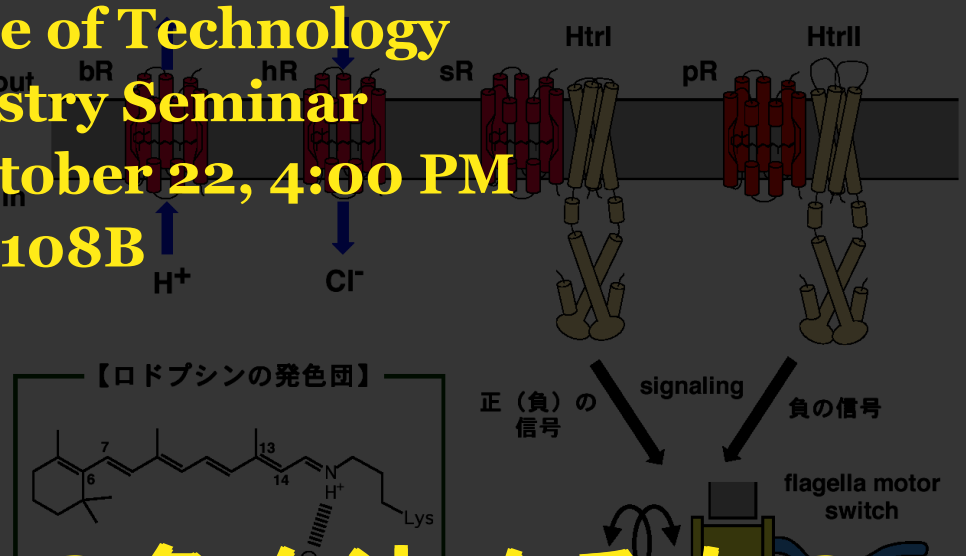




Nagoya Institute of Technology  
 Physical Chemistry Seminar  
 Wednesday, October 22, 4:00 PM  
 1st Bldg, Room 108B

【光エネルギー変換】

【光情報変換】



# 古細菌ロドプシンの色を決めるもの

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(北海道大学大学院薬学研究科)

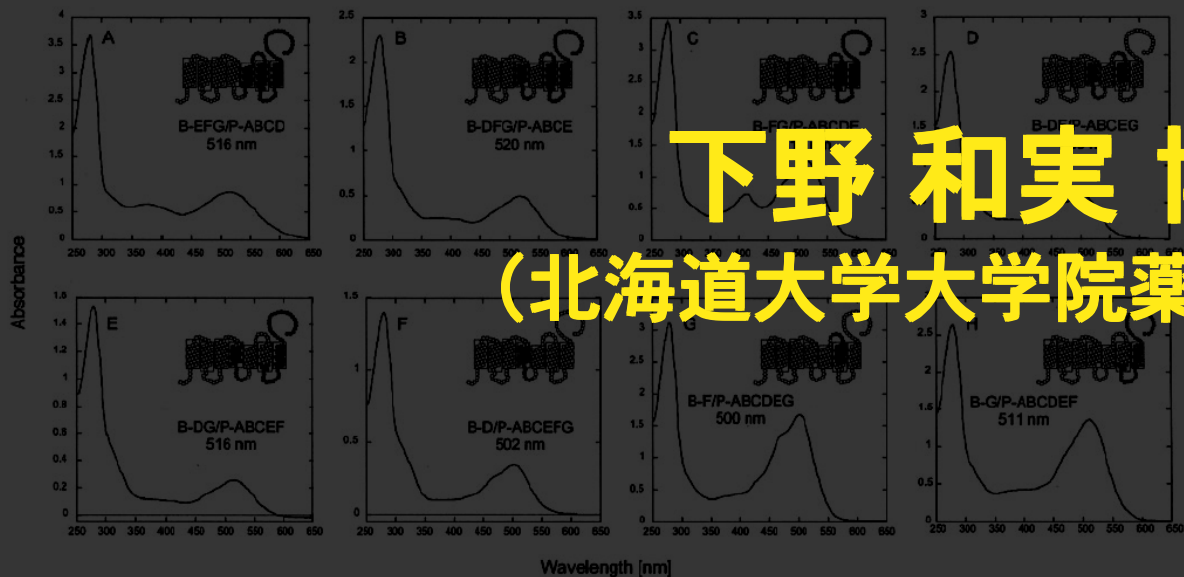


FIG. 4. UV-Vis spectra of various chimeric proteins. Opsin types and absorption maxima are indicated in each panel. The pigment structures are shown in each panel. The residues from bR are shown as filled circles, and those from ppR as open circles. The experimental conditions and the notations are the same as described in the legend for Fig. 2.



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FIG. 6. Specific amino acid residues for determining factors at helix D, E, and G. The structures of bR (Protein Data Bank (PDB) code, 1C3W) and ppR (PDB code, 1C3V) are shown, respectively. The structural alignment is done for the all atoms of the helix D and Lys-205 for ppR and Lys-216 for bR.