

Lampiran 1. Routine Program Simulasi

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%===== Membuat jendela halaman utama ======%

figure1 = figure('PaperSize',[1 1],'PaperUnits','normalized',...
    'NumberTitle','off',...
    'Name','Halaman Utama',...
    'menubar','none',...
    'Color',[0.8 0.8 0.8]);
panel_utama=uipanel('parent',figure1,'Title','Thesis','FontSize',12,...
    'BackgroundColor',[0.8 0.8 0.8], ...
    'FontWeight','bold','Fontangle',...
    'Italic','shadowcolor',[1 1 0],...
    'ForegroundColor',[1 1 0],'Position',[.02 .02 0.96 0.96]);
axes1=axes('Position',[0.4326 0.2982 0.1738 0.1])
logo_TI=imread('logo TI.jpg');
image(logo_TI);
axis off;

teks1=annotation(figure1,'textbox',[0.4424 0.505 0.1553 0.09955],...
    'String',{'Oleh :','Triyono Budi Santoso','0806422813'},...
    'HorizontalAlignment','center',...
    'FontWeight','bold',...
    'FontSize',12,...
    'FitBoxToText','on',...
    'FontName','Times New Roman',...
    'LineStyle','none');

teks2=annotation(figure1,'textbox',[0.3041 0.1651 0.4393 0.09007],...
    'String',{'DEPARTEMEN TEKNIK INDUSTRI','PROGRAM PASCA
SARJANA','UNIVERSITAS INDONESIA','2010'},...
    'HorizontalAlignment','center',...
    'FontWeight','bold',...
    'FontSize',14,...
    'FontName','Times New Roman',...
    'FitBoxToText','on',...
    'LineStyle','none');

teks3=annotation(figure1,'textbox',[0.2827 0.80 0.4561 0.1085],...
    'String',{' ANALISIS KUALITAS REDAMAN SERAT OPTIK UNTUK',
'MENINGKATKAN KINERJA SISTEM TELEKOMUNIKASI',
'DENGAN MENGGUNAKAN DOE'},...
    'HorizontalAlignment','center',...
    'FontWeight','bold',...
    'FontSize',14,...
    'FitBoxToText','on',...
    'FontName','Times New Roman',...
    'LineStyle','none',...
    'EdgeColor',[0 0 1]);
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align_teks=[teks1 teks2 teks3]
align(align_teks,'distribute','centre')
tombol_simulasi=uicontrol(panel_utama,'style','pushbutton','string','Simulasi','unit
s','normalized','callback','jendela_doe',...
'Position',[0.87 0.01 0.1 0.05])

close;
%jendela pertanyaan
fig_pertanyaan_jum_per_bending=figure('Units','normalized',...
'Position',[0.3 0.5 0.4 0.1],...
'Color',[1 1 0],...
'HandleVisibility','callback',...
'NumberTitle','off',...
'Name','Masukkan Jumlah Pengukuran',...
'Menubar','none');

%Text pertanyaan
jum_per_splicing=annotation(fig_pertanyaan_jum_per_bending,'textbox',[0.275
0.6028 0.1852 0.06048],...
'String',{'Jumlah Pengukuran =      kali'},...
'LineStyle','none',...
'FitBoxtoText','on');

%Edit jumlah percobaan
edit_jumlah_pertanyaan = uicontrol(fig_pertanyaan_jum_per_bending,'Style',
'edit','Unit','normalized','Position',[0.55 0.35 0.06 0.26048]);
%Tombol Lanjutkan
tombol_lanjutkan =
uicontrol(fig_pertanyaan_jum_per_bending,'style','pushbutton',...
'string','Lanjutkan','Unit','normalized','Position', [0.8 0.15 0.15 0.26048],
'Callback','data_percobaan_faktor_tunggal_bending');

%=====Mengumpulkan data percobaan pengukuran
splicing=====
%Membaca input jumlah percobaan
edit_jumlah_percobaan=get(edit_jumlah_pertanyaan,'string')
jumlah=str2num(edit_jumlah_percobaan)
close;
fig_percobaan = figure('PaperSize',[1 1],'PaperUnits','normalized',...
'NumberTitle','off',...
'Name','Data Percobaan Splicing','resize','off', ...
'menubar','none',...
'Color',[0.8 0.8 0.8]);
uipanel('parent',figure1,'Title','Kombinasi Faktor (F) & Level
(L)','FontSize',12,'BackgroundColor',[0.8 0.8 0.8], ...
'FontWeight','bold','Fontangle','Italic','shadowcolor',[1 1 0],...
'ForegroundColor',[1 1 0],'Position',[.02 .02 0.96 0.96]);
matrix_percobaan = zeros (jumlah,6); % 6 adalah 2 x 3 level

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cnames = {'F1L1-F2L1','F1L1-F2L2','F1L1-F2L3','F1L2-F2L1','F1L2-
F2L2','F1L2-F2L3'};

%===== Menganalisis data percobaan pengukuran bending
=====
h = waitbar(0,'Sedang Memproses, Silakan Tunggu....');
for i=1:4000,
waitbar(i/4000)
end
close(h)
%data_retrieve=get(tabel_percobaan,'data')
data_retrieve=xlsread('data_bending_final.xlsx');
close
%Menguji normalitas
input('1. Analisis Normalitas')
baca_data_bending =data_retrieve;
figure
plot_bending1=subplot(3,1,1);
normplot(baca_data_bending(:,1));
set(get(plot_bending1,'Xlabel'),'string','Brand 1')
title('')
plot_bending2=subplot(3,1,2);
normplot(baca_data_bending(:,2));
set(get(plot_bending2,'Xlabel'),'string','Brand 2')
title('')
plot_bending3=subplot(3,1,3);
normplot(baca_data_bending(:,3));
set(get(plot_bending3,'Xlabel'),'string','Brand 3')
title('')
set(gcf,'numbertitle','on','name','Plot normalitas data redaman bending untuk tiap
brand FO')
pause
close all;
%Menguji homogenitas varians
input('2. Analisis Homogenitas Varians')
data_retrieve_homogen = data_retrieve (:);

uji_homogen_bartlett=uji_varian (data_retrieve_homogen,group_respon);% uji
Bartlett
close;
set(gcf,'numbertitle','on','name','Uji Varians metode Bartlett');
uji_homogen_levene=uji_varian
(data_retrieve_homogen,group_respon,'on','robust'); % uji Levene
close;
set(gcf,'numbertitle','on','name','Uji Varians metode Levene');
pause
close all;

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%Menganalisis data pengukuran dengan menggunakan ANOVA 1 Arah
input('3. Analisis Varians Satu Arah')
data_matrix=data_retrieve;

[p,tbl,stats] = anova_satu_arah_bending(data_matrix);
set(get(gca,'XLabel'),'String','Brand FO')
set(get(gca,'YLabel'),'String','Nilai Rata-rata')
figure
[c,m] = multcompare(stats); %untuk menguji perbedaan mean
set(gcf,'numbertitle','on','name','Perbandingan rata-rata redaman bending untuk
tiap brand FO')
set(get(gca,'XLabel'),'String','Redaman')
set(get(gca,'YLabel'),'String','Brand FO')
set(get(gca,'Title'),'String','')

%===== Menganalisis data percobaan pengukuran patching
=====
h = waitbar(0,'Sedang Memproses, Silakan Tunggu....');
for i=1:4000,
waitbar(i/4000)
end
close(h)
%data_retrieve=get(tabel_percobaan,'data')
data_retrieve=xlsread('data_patching_final.xlsx');
close;
%Menguji normalitas
input('1. Analisis Normalitas')
baca_data_patching =data_retrieve;
figure
plot_patching1=subplot(2,2,1);
normplot(baca_data_patching(:,1));
set(get(plot_patching1,'Xlabel'),'string','Outdoor')
title('ODF Tipe 1','Color','blue','FontWeight','bold')
plot_patching2=subplot(2,2,2);
normplot(baca_data_patching(:,3));
set(get(plot_patching2,'Xlabel'),'string','Outdoor')
title('ODF Tipe 2','Color','blue','FontWeight','bold')
plot_patching3=subplot(2,2,3);
normplot(baca_data_patching(:,2));
set(get(plot_patching3,'Xlabel'),'string','Indoor')
title('')
plot_patching4=subplot(2,2,4);
normplot(baca_data_patching(:,4));
set(get(plot_patching4,'Xlabel'),'string','Indoor')
title('')
set(gcf,'numbertitle','on','name','Plot normalitas data redaman patching untuk tiap
jenis ODF dan lokasi penempatan')
pause
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close all;

data_retrieve1=data_retrieve(:,1:2);
data_matrix1=data_retrieve1(:);
data_retrieve2=data_retrieve(:,3:4);
data_matrix2=data_retrieve2(:);
data_matrix=[data_matrix1 data_matrix2];

%Menganalisis data pengukuran dengan menggunakan ANOVA 2 Arah
input('3. Analisis Varians Dua Arah')
reps=jumlah;
anova_dua_arah_patching(data_matrix,jumlah);
set(gcf,'numbertitle','on')

%Plotting Interaksi
n=jumlah;
a=2;
b=2;
t=a*b*n;
respon1=[data_retrieve(:,1); data_retrieve(:,3)];
respon2=[data_retrieve(:,2); data_retrieve(:,4)];
respon =[respon1;respon2];
group1=zeros(t,1);
m=0;
for i=1:b
    u=n*a*m+1;
    o=n*a*i;
    group1(u:o,1)=m+1;
    m=i;
end
group2=zeros(t,1);
m=0;
for i=1:b
    k=n*m+1;
    l=n*i;
    group2(k:l,1)=m+1;
    m=i;
end
h=0;
for i=1:b
    group2(n*a*h+1:n*a*i,1)=group2(1:n*a,1);
    h=i;
end
group=[group1 group2];
figure1 = figure('PaperSize',[1 1],'PaperUnits','normalized',...
    'NumberTitle','on',...
    'Name','Grafik Interaksi',...
    'menubar','none',...

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'Color',[0.8 0.8 0.8]);
interactionplot(respon,group,'varnames',{'Lokasi ODF','Jenis ODF'})
%Plotting Pengaruh Utama
figure1 = figure('PaperSize',[1 1],'PaperUnits','normalized',...
    'NumberTitle','on',...
    'Name','Pengaruh Utama',...
    'menubar','none',...
    'Color',[0.8 0.8 0.8]);
maineffectsplot(respon,{group1, group2}, 'varnames',{'Lokasi ODF','Jenis ODF'})
set(get(gca,'YLabel'), 'String', 'Redaman')

% fungsi ANOVA %
function [p,anovatab,stats] = anova1(x,group,displayopt,extra)
classical = 1;
nargs = nargin;
if (nargin > 0 && strcmp(x,'kruskalwallis'))
    % Fungsi kruskalwallis adjust input
    classical = 0;
    if (nargin >= 2), x = group; group = []; end
    if (nargin >= 3), group = displayopt; displayopt = []; end
    if (nargin >= 4), displayopt = extra; end
    nargs = nargs-1;
end
error(nargchk(1,3,nargs,'struct'));

if (nargs < 2), group = []; end
if (nargs < 3), displayopt = 'on'; end

willdisplay = ~(strcmp(displayopt,'nodisplay') | strcmp(displayopt,'n') ...
    | strcmp(displayopt,'off'));

% Merubah group ke cell array dari karakter array dan membuatnya dalam 1
% kolom
if (ischar(group) && ~isempty(group)), group = cellstr(group); end
if (size(group, 1) == 1), group = group'; end

% Mengambil ukuran group dan mean/rata-rata
k = find(groupnum == j);
lk = length(k);
countx(j) = lk;
xm(j) = mean(xr(k));      % kolom mean

if (willdisplay)          % boxplot
    M(1:lk,j) = xorig(k);
end
end

gm = mean(xr);           % grand mean

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df1 = sum(countx>0) - 1;           % dof kolom
df2 = lx - df1 - 1;               % dof error
xc = xm - gm;
xc(countx==0) = 0;
RSS = dot(countx, xc.^2);         % Regressi Sum of Squares
else

[r,c] = size(x);
lx = r * c;
if (classical)
    xr = x;
    mu = mean(xr(:));
    xr = xr - mu;
else
    [xr,tieadj] = tiedrank(x(:));
    xr = reshape(xr, size(x));
end
countx = repmat(r, 1, c);
xorig = x;                      % save boxplot
xm = mean(xr);                  % kolom mean/rata-rata
gm = mean(xm);                  % grand mean
df1 = c-1;                       % dof kolom
df2 = c*(r-1);                  % dof error
RSS = r*(xm - gm)*(xm-gm)';     % Regressi Sum of Squares
end

TSS = (xr(:) - gm)'*(xr(:) - gm); % Total Sum of Squares
SSE = TSS - RSS;                 % Error Sum of Squares

if (df2 > 0)
    mse = SSE/df2;
else
    mse = NaN;
end

if (classical)
    if (SSE~=0)
        F = (RSS/df1) / mse;
        p = 1 - fcdf(F,df1,df2);   % Probability dari F
    elseif (RSS==0)
        F = NaN;
        p = NaN;
    else
        F = Inf;
        p = 0;
    end
else
    F = (12 * RSS) / (lx * (lx+1));
end

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if (tieadj > 0)
    F = F / (1 - 2 * tieadj/(lx^3-lx));
end
p = 1 - chi2cdf(F, df1);
end

Table=zeros(3,5);          %Format Tabel ANOVA
Table(:,1)=[ RSS SSE TSS]';
Table(:,2)=[df1 df2 df1+df2]';
Table(:,3)=[ RSS/df1 mse Inf ]';
Table(:,4)=[ F Inf Inf ]';
Table(:,5)=[ p Inf Inf ]';

colheads = ['Source      ';'   SS  ';'   df';...
            '   MS  ';'     F  ','  Prob>F '];
if (~classical)
    colheads(5,:) = ' Chi-sq ';
    colheads(6,:) = ' Prob>Chi-sq';
end
rowheads = ['Brand FO  ';'Error      ';'Total      '];
if (grouped)
    rowheads(1,:)= 'Groups  ';
end

% Tabel cell array
atab = num2cell(Table);
for i=1:size(atab,1)
    for j=1:size(atab,2)
        if (isinf(atab{i,j}))
            atab{i,j} = [];
        end
    end
end
atab = [cellstr(strjust(rowheads, 'left')), atab];
atab = [cellstr(strjust(colheads, 'left'))'; atab];
if (nargout > 1)
    anovatab = atab;
end

% Multicompare
if (nargout > 2)
    if (length(gnames) > 0)
        stats.gnames = gnames;
    else
        stats.gnames = strjust(num2str((1:length(xm))), 'left');
    end
    stats.n = countx;
    if (classical)

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stats.source = 'anova1';
stats.means = xm + mu;
stats.df = df2;
stats.s = sqrt(mse);
else
    stats.source = 'kruskalwallis';
    stats.meanranks = xm;
    stats.sumt = 2 * tieadj;
end
end

if (~willdisplay), return; end

digits = [-1 -1 0 -1 2 4];
if (classical)
    wtitle = 'Karakteristik redaman bending untuk tiap brand FO';
    ttitle = 'Tabel ANOVA';
else
    wtitle = 'ANOVA Kruskal-Wallis';
    ttitle = 'Tabel ANOVA Kruskal-Wallis';
end
tblfig = statdisptable(atab, wtitle, ttitle, ", digits);
set(tblfig,'tag','table','NumberTitle','on');

f1 = figure('pos',get(gcf,'pos') + [0,-200,0,0], 'tag','boxplot','Name','Diagram Box
Plot','NumberTitle','on');
ax = axes('Parent',f1);
if (~grouped)
    boxplot(ax,xorig,'notch','on');
else
    boxplot(ax,M,'notch','on');
    h = get(ax,'XLabel');
    set(h,'String','Group Number');
end

colmean = mean(Y,1);      % rata-rata kolom
rowmean = mean(Y,2);      % rata-rata baris
gm = mean(colmean);       % rata-rata total
df1 = c-1;                 % dof kolom
df2 = m-1;                 % dof baris
if reps == 1,
    edf = (c-1)*(r-1);     % dof error tanpa replikasi
else
    edf = (c*m*(reps-1));   % dof error dengan replikasi
    idf = (c-1)*(m-1);      % dof interaksi
end
CSS = m*reps*(colmean - gm)*(colmean-gm)';           % Sum of Squares kolom
RSS = c*reps*(rowmean - gm)*(rowmean-gm)';           % Sum of Squares baris

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correction = (c*m*reps)*gm^2;
TSS = sum(sum(X .* X)) - correction; % Sum of Squares total
ISS = reps*sum(sum(Y .* Y)) - correction - CSS - RSS; % Sum of Squares
interaksi
if reps == 1,
    SSE = ISS;
else
    SSE = TSS - CSS - RSS - ISS; % Sum of Squares error
end

ip = NaN;
if (SSE~=0)
    MSE = SSE/edf;
    colf = (CSS/df1) / MSE;
    rowf = (RSS/df2) / MSE;
    colp = 1 - fcdf(colf,df1,edf); % Probability F untuk rata-rata kolom
    rowp = 1 - fcdf(rowf,df2,edf); % Probability F untuk rata-rata baris
    p = [colp rowp];

    if (reps > 1),
        intf = (ISS/df)/MSE;
        ip = 1 - fcdf(intf,df,edf);
        p = [p ip];
    end
else
    if (edf> 0)
        MSE = 0;
    else
        MSE = NaN;
    end
    if CSS==0, % Tidak ada variansi antar kolom
        colf = NaN;
        colp = NaN;
    else % Ada variansi antar kolom
        colf = Inf;
        colp = 0;
    end

    if RSS==0, % Tidak ada variansi antar baris
        rowf = NaN;
        rowp = NaN;
    else % Ada variansi antar baris
        rowf = Inf;
        rowp = 0;
    end

    p = [colp rowp];

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```

if (reps>1) && (ISS==0) % Replikasi tapi tidak ada interaksi
    intf = NaN;
    p = [p NaN];
elseif (reps>1)      % Replikasi dengan interaksi
    intf = Inf;
    p = [p 0];
end

if (reps > 1),
    Table{6,6} = []; %Format tabel ANOVA Table dengan interaksi
    Table(2:6,1)={Jenis Splicer';'Brand FO';'Interaction';'Error';'Total'};
    Table(2:6,2)={CSS; RSS; ISS; SSE; TSS};
    Table(2:6,3)={df1; df2; idf; edf; r*c-1};
    Table(2:5,4)={CSS/df1; RSS/df2; ISS/idf; SSE/edf;};
    Table(2:4,5)={colf; rowf; intf};
else
    Table{5,6} = []; %Format tabel ANOVA tanpa interaksi
    Table(2:5,1)={Jenis Splicer';'Brand FO';'Error';'Total'};
    Table(2:5,2)={CSS; RSS; SSE; TSS};
    Table(2:5,3)={df1; df2; edf; r*c-1};
    Table(2:4,4)={CSS/df1; RSS/df2; SSE/edf;};
    Table(2:3,5)={colf; rowf};
end
Table(1,1:6) = {'Source' 'SS' 'df' 'MS' 'F' 'Prob>F'};
Table(2:(1+length(p)),6) = num2cell(p);

if (isequal(displayopt, 'on'))
    digits = [-1 -1 0 -1 2 4];
    statdisptable(Table, 'Karakteristik redaman splicing dari tiap splicer untuk tiap
brand FO', 'Tabel ANOVA', ", digits);
end

if (nargout > 2)
    stats.source = 'anova2';
    stats.sigmasq = MSE;
    stats.colmeans = colmean; % rata-rata kolom
    stats.coln = m*reps;      % n untuk estimasi rata-rata kolom
    stats.rowmeans = rowmean;
    stats.rown = c*reps;
    stats.inter = (reps>1);
    stats.pval = ip;          % p-value interaksi
    stats.df = edf;
end

function [p,Table,stats] = anova2(X,reps,displayopt)
if (nargin<3), displayopt = 'on'; end

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```

if (nargin < 1)
    error('stats:anova2:TooFewInputs','At least one input is required.');
end
if (any(isnan(X(:))))
    error('stats:anova2:DataNotBalanced',...
        'NaN values in input not allowed. Use anovan instead.');
end
[r,c] = size(X);
if nargin == 1,
    reps = 1;
    m=r;
    Y = X;
elseif reps == 1
    m=r;
    Y = X;
else
    m = r/reps;
    if (floor(m) ~= r/reps),
        error('stats:anova2:BadSize',...
            'The number of rows must be a multiple of reps.');
    end
    Y = zeros(m,c);
    for i=1:m,
        j = (i-1)*reps;
        Y(i,:) = mean(X(j+1:j+reps,:));
    end
end
colmean = mean(Y,1);      % rata-rata kolom
rowmean = mean(Y,2>');   % rata-rata baris
gm = mean(colmean);       % rata-rata total
df1 = c-1;                 % dof kolom
df2 = m-1;                 % dof baris
if reps == 1,
    edf = (c-1)*(r-1);     % dof error tanpa replikasi
else
    edf = (c*m*(reps-1));  % dof error dengan replikasi
    idf = (c-1)*(m-1);     % dof interaksi
end
CSS = m*reps*(colmean - gm)*(colmean-gm)';           % Sum of Squares kolom
RSS = c*reps*(rowmean - gm)*(rowmean-gm)';           % Sum of Squares baris
correction = (c*m*reps)*gm^2;
TSS = sum(sum(X .* X)) - correction;                  % Sum of Squares total
ISS = reps*sum(sum(Y .* Y)) - correction - CSS - RSS; % Sum of Squares
interaksi
if reps == 1,
    SSE = ISS;
else
    SSE = TSS - CSS - RSS - ISS;                      % Sum of Squares error

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end

ip = NaN;
if (SSE~=0)
    MSE = SSE/edf;
    colf = (CSS/df1) / MSE;
    rowf = (RSS/df2) / MSE;
    colp = 1 - fcdf(colf,df1,edf); % Probability F untuk rata-rata kolom
    rowp = 1 - fcdf(rowf,df2,edf); % Probability F untuk rata-rata baris
    p = [colp rowp];

if (reps > 1),
    intf = (ISS/idf)/MSE;
    ip = 1 - fcdf(intf,idf,edf);
    p = [p ip];
end

else
    if (edf > 0)
        MSE = 0;
    else
        MSE = NaN;
    end
    if CSS==0,      % Tidak ada variansi antar kolom
        colf = NaN;
        colp = NaN;
    else          % Ada variansi antar kolom
        colf = Inf;
        colp = 0;
    end

    if RSS==0,      % Tidak ada variansi antar baris
        rowf = NaN;
        rowp = NaN;
    else          % Ada variansi antar baris
        rowf = Inf;
        rowp = 0;
    end

p = [colp rowp];

if (reps>1) && (ISS==0) % Replikasi tapi tidak ada interaksi
    intf = NaN;
    p = [p NaN];
elseif (reps>1)      % Replikasi dengan interaksi
    intf = Inf;
    p = [p 0];
end

```

```

end

if (reps > 1),
Table{6,6} = []; %Format tabel ANOVA Table dengan interaksi
Table(2:6,1)={'Jenis ODF'; 'Lokasi'; 'Interaction'; 'Error'; 'Total'};
Table(2:6,2)={CSS; RSS; ISS; SSE; TSS};
Table(2:6,3)={df1; df2; idf; edf; r*c-1};
Table(2:5,4)={CSS/df1; RSS/df2; ISS/idf; SSE/edf;};
Table(2:4,5)={colf; rowf; intf};
else
Table{5,6} = []; %Format tabel ANOVA tanpa interaksi
Table(2:5,1)={'Jenis ODF'; 'Lokasi'; 'Error'; 'Total'};
Table(2:5,2)={CSS; RSS; SSE; TSS};
Table(2:5,3)={df1; df2; edf; r*c-1};
Table(2:4,4)={CSS/df1; RSS/df2; SSE/edf;};
Table(2:3,5)={colf; rowf};
end

Table(1,1:6) = {'Source' 'SS' 'df' 'MS' 'F' 'Prob>F'};
Table(2:(1+length(p)),6) = num2cell(p);

if (isequal(displayopt, 'on'))
    digits = [-1 -1 0 -1 2 4];
    statdisptable(Table, 'Karakteristik redaman patching dari tiap ODF untuk tiap lokasi penempatan', 'Tabel ANOVA', "", digits);
end

if (nargout > 2)
    stats.source = 'anova2';
    stats.sigmasq = MSE;
    stats.colmeans = colmean; % rata-rata kolom
    stats.coln = m*reps;      % n untuk estimasi rata-rata kolom
    stats.rowmeans = rowmean;
    stats.rown = c*reps;
    stats.inter = (reps>1);
    stats.pval = ip;          % p-value interaksi
    stats.df = edf;
end

%===== Tes data bending =====%
baca_data_bending = xlsread('data_bending_final.xlsx');
figure
subplot(3,1,1);
normplot(baca_data_bending(:,1));
subplot(3,1,2);
normplot(baca_data_bending(:,2));
subplot(3,1,3);
normplot(baca_data_bending(:,3));

```

```

set(gcf,'numbertitle','on','name','Plot Normalitas Data Bending')

%===== Tes data splicing =====%
baca_data_splicing =xlsread('data_splicing_final.xlsx');
subplot(3,2,1);
normplot(baca_data_splicing(:,1));
subplot(3,2,2);
normplot(baca_data_splicing(:,4));
subplot(3,2,3);
normplot(baca_data_splicing(:,2));
subplot(3,2,4);
normplot(baca_data_splicing(:,5));
subplot(3,2,5);
normplot(baca_data_splicing(:,3));
subplot(3,2,6);
normplot(baca_data_splicing(:,6));
set(gcf,'numbertitle','on','name','Plot Normalitas Data Splicing')

%===== Tes data patching =====%
baca_data_patchting =xlsread('data_patchting_final.xlsx');
figure
subplot(2,2,1);
normplot(baca_data_patchting(:,1));
subplot(2,2,2);
normplot(baca_data_patchting(:,3));
subplot(2,2,3);
normplot(baca_data_splicing(:,2));
subplot(2,2,4);
normplot(baca_data_patchting(:,4));
set(gcf,'numbertitle','on','name','Plot Normalitas Data Patching')

% === uji varian ===%
function [p,stats] = vartestn(x,group,displayopt,testtype)

error(nargchk(1,4,nargin,'struct'));

if nargin < 2, group = []; end
if nargin < 3 || isempty(displayopt)
    displayopt = 'on';
elseif ~isequal(displayopt,'on') && ~isequal(displayopt,'off')
    error('stats:vartestn:BadDisplayOpt',...
        'DISPLAYOPT must be "on" or "off"!');

end
dodisplay = isequal(displayopt,'on');

if nargin < 4 || isempty(testtype)
    dorobust = false;

```

```

else
    if ischar(testtype)
        i = find(strncmpi(testtype, {'robust';'classical'},length(testtype)));
    else
        i = [];
    end
    if isempty(i)
        error('stats:vartestn:BadTestType',...
            'TESTTYPE must be one of the strings "robust" or "classical".');
    end
    dorobust = (i==1);
end

% Error jika data tidak ada
if isempty(x)
    error('stats:vartestn>NoData','X must not be empty');
end

% Merubah group ke array
if (ischar(group) && ~isempty(group)), group = cellstr(group); end
if (size(group, 1) == 1), group = group'; end

% jika x matrix, ubah ke bentuk vektor.
[n,m] = size(x);
if isempty(group)
    x = x(:);
    group = reshape(repmat((1:m), n, 1), n*m, 1);
elseif isvector(x)
    if numel(group) ~= n
        error('stats:vartestn:InputSizeMismatch',...
            'X and GROUP must have the same length.');
    end
else
    error('stats:vartestn:BadGroup',...
        'Cannot specify a GROUP argument unless X is a vector.');
end

% NaN
t = isnan(x);
if any(t)
    x(t) = [];
    group(t,:) = [];
end

% Menghitung summary group
[igroup,gname] = grp2idx(group);
[gmean,gsem,gcount] = grpstats(x,igroup);
df = gcount-1;           % group degrees of freedom

```

```

gvar = gcount .* gsem.^2;      % group variances
sumdf = sum(df);
if sumdf>0
    vp = sum(df.*gvar) / sumdf; % pooled variance
else
    vp = NaN;
end
k = length(df);

if dorobust
    % Levene's test

    spgroups = find(gcount<2);
    t = ~ismember(igroup,spgroups);
    xc = x(t) - gmean(igroup(t));
    ngroups = length(gcount) - length(spgroups);

    % ANOVA
    if ngroups>1
        [p,atab] = anova1(xc.^2, igroup,'off');
        B = atab{2,5};           % F statistik
        Bdf = [atab{2,3}, atab{3,3}]; % d.f.
    else
        p = NaN;
        B = NaN;
        Bdf = [0, length(xc)-ngroups];
    end
    Bdftable = sprintf("%d, %d",Bdf(1),Bdf(2));
    testname = 'Levene''s statistic';
    statname = 'fstat';
else
    % Bartlett's test
    Bdf = max(0, sum(df>0)-1);
    t = df>0;
    if Bdf>0 && sumdf>0
        B = log(vp) * sum(df) - sum(df(t).*log(gvar(t)));
        C = 1 + (sum(1./df(t)) - 1/sum(df))/(3*Bdf);
        B = B/C;
    else
        B = NaN;
    end

    p = 1 - chi2cdf(B,Bdf);
    Bdftable = Bdf;
    testname = 'Bartlett''s statistic';
    statname = 'chisqstat';
end

```

```

if dodisplay
    Table = cell(k+6,4);
    Table(1,:) = {'Group' 'Count' 'Mean' 'Std Dev'};
    Table(2:k+1,1) = gname;
    Table(2:k+1,2) = num2cell(gcount);
    Table(2:k+1,3) = num2cell(gmean);
    Table(2:k+1,4) = num2cell(sqrt(gvar));

    Table{k+2,1} = 'Pooled';
    Table{k+2,2} = sum(gcount);
    Table{k+2,3} = sum(gcount.*gmean) / sum(gcount);
    Table{k+2,4} = sqrt(vp);

    Table{k+3,1} = '';
    Table{k+4,1} = testname;
    Table{k+4,2} = B;
    Table{k+5,1} = 'Degrees of freedom';
    Table{k+5,2} = Bdftable;
    Table{k+6,1} = 'p-value';
    Table{k+6,2} = p;

tblfig = statdisptable(Table, 'Variance Test', 'Tabel Uji Varians');
set(tblfig,'tag','table');
end

% Membuat struktur output
if (nargout > 1)
    stats = struct(statname,B,'df',Bdf);
end

if ~dodisplay
    return;
end

% Membuat boxplot
f1 = figure('pos',get(gcf,'pos') + [0,-200,0,0], 'tag', 'boxplot');
ax = axes('Parent',f1);
boxplot(ax,x,group);

% ===== Fungsi Multicompare=====%
function [comparison,means,h,gnames] = multcompare(stats,varargin)
if (nargin < 1)
    error('stats:multcompare:TooFewInputs',...
        'At least one argument is required.');
end
if isempty(varargin) || (~isempty(varargin{1}) && ischar(varargin{1}))
    okargs = {'alpha' 'display' 'ctype' 'estimate' 'dimension'};

```

```

defaults = {0.05 'on' " " []};
[eid,emsg,alpha,displayopt,ctype,estimate,dim] = ...
    statgetargs(okargs,defaults,varargin{:});
if ~isempty(emsg)
    error(sprintf('stats:multcompare:%s',eid),emsg);
end
else

    if (nargin >= 2) && ~isempty(varargin{1})
        alpha = varargin{1};
    else
        alpha = 0.05;
    end
    if (nargin >= 3) && ~isempty(varargin{2})
        displayopt = varargin{2};
    else
        displayopt = 'on';
    end
    if (nargin>=4) && ~isempty(varargin{3})
        ctype = varargin{3};
    else
        ctype = "";
    end
    if (nargin>=5) && ~isempty(varargin{4})
        estimate = varargin{4};
    else
        estimate = "";
    end
    if (nargin>=6) && ~isempty(varargin{5})
        dim = varargin{5};
    else
        dim = [];
    end
end

if (~isstruct(stats))
    error('stats:multcompare:BadStats','STATS must be a structure.');
end
if (~isfield(stats,'source'))
    error('stats:multcompare:BadStats',...
        'The first argument is not a STATS output structure from ANOVA1!');

if (length(alpha)~=1 || ~isfinite(alpha) || alpha<=0 || alpha>=1)
    error('stats:multcompare:BadAlpha',...
        'ALPHA must be a scalar between 0 and 1, exclusive.');
end
if ~(isequal(displayopt, 'on') || isequal(displayopt, 'off'))
    error('stats:multcompare:BadDisplayOpt',...

```

```

'DISPLAYOPT must be "on" or "off".');
end
dodisp = isequal(displayopt, 'on');

source = stats.source;

switch(source)
case 'anova1'
    mname = 'means';
    gmeans = stats.means();
    gnames = stats.gnames;
    n = stats.n();
    df = stats.df;
    s = stats.s;
    ng = sum(n>0);
    if (df< 1)
        error('stats:multcompare:NotEnoughData',...
            'Cannot compare means with 0 degrees of freedom for error.');
    end

% Tukey-Kramer
if (isempty(ctype)), ctype = 'tukey-kramer'; end
crit = getcrit(ctype, alpha, df, ng);

gcov = diag((s^2)./n);

case 'anova2'
    docols = logical(1);
    if (~isempty(estimate))
        if ~(~isequal(estimate,'row') || isequal(estimate,'column')...
            || isequal(estimate,'r') || isequal(estimate,'c'))
            error('stats:multcompare:BadEstimate',...
                'ESTIMATE must be "column" or "row".');
        end
        docols = isequal(estimate, 'column') || isequal(estimate, 'c');
    end
    if (docols)
        gmeans = stats.colmeans();
        n = stats.coln();
        mname = 'column means';
    else
        gmeans = stats.rowmeans();
        n = stats.rown();
        mname = 'row means';
    end
    ng = length(gmeans);
    sigma = sqrt(stats.sigmasq);
    gnames = strjust(num2str((1:ng)'), 'left');

```

```

df = stats.df;
if (df < 1)
    error('stats:multcompare:NotEnoughData',...
        'Cannot compare means with 0 degrees of freedom for error.');
end

% Tukey-Kramer
if (isempty(ctype)), ctype = 'tukey-kramer'; end
crit = getcrit(ctype, alpha, df, ng);

gcov = ((sigma^2)/n) * eye(ng);

if (stats.inter && dodisp)
    if (stats.pval < alpha)
        disp(sprintf(...

['Note: Your model includes an interaction term that is significant\n',...
    'at the level you specified. Testing main effects under these\n',...
    'conditions is questionable.']));
    else
        disp(sprintf(...

['Note: Your model includes an interaction term. A test of main\n',...
    'effects can be difficult to interpret when the model includes\n',...
    'interactions.']));

    end
end

case 'anovan'
    mname = 'population marginal means';

    try
        vnested = stats.vnested;
    catch
        vnested = [];
    end
    if any(vnested(:))
        error('stats:multcompare:NoNesting',...
            'MULTCOMPARE does not support models with nested factors.');
    end

    if ~isempty(stats.ems)
        warning('stats:multcompare:IgnoringRandomEffects',...
            'Ignoring random effects (all effects treated as fixed.)')
    end

nvars = length(stats.nlevels);
P0 = stats.nullproject;

```

```

if isempty(dim)
    dim = find(stats.nlevels>1,1);
end
dim = dim(:);
if isempty(dim) || any(dim<1 | dim>nvars | dim~=round(dim))
    error('stats:multcompare:BadDim',...
        'Values of DIM must be integers between 1 and %d.',nvars);
end
dim = sort(dim);
dim(diff(dim)==0) = [];
if any(stats.nlevels(dim)<2)
    error('stats:multcompare:BadDim',...
        'DIM must specify only categorical factors with 2 or more degrees of
freedom.');
end

try
    continuous = stats.continuous;
catch
    continuous = zeros(nvars,1);
end
ffdesign = fullfact(stats.nlevels(dim));
nrows = size(ffdesign,1);

dums = cell(nvars, 1);
for j=1:length(dim)
    dj = dim(j);
    dums{dj} = idummy(ffdesign(:,j),3);
end

for j=1:nvars
    if isempty(dums{j});
        if continuous(j)
            dums{j} = repmat(stats.vmeans(j),nrows,1);
        else
            nlev = stats.nlevels(j);
            dums{j} = repmat(1/nlev,nrows,nlev);
        end
    end
end

termcols = stats.termcols();
termstart = cumsum([1; termcols]);
terms = [zeros(1,nvars); stats.terms];
ncols = sum(termcols);
x = zeros(size(ffdesign,1), ncols);
x(:,1) = 1;
for j=1:length(termcols)

```

```

tm = terms(j,:);
t0 = termstart(j);
t1 = termstart(j) + termcols(j) - 1;
if all(tm==0)
    x(:,t0:t1) = 1;
else
    x0 = [];
    for k=nvars:-1:1
        if tm(k)
            x0 = termcross(x0,dums{k});
        end
    end
    x(:,t0:t1) = x0;
end
end

% Estimasi error
gmeans = x * stats.coeffs;
xproj = (x*P0)';
tmp = stats.Rtr \ xproj;
if (stats.dfe == 0)
    mse = NaN;
else
    mse = max(stats.mse,0);
end
gcov = mse * tmp' * tmp;

Xrows = stats.rowbasis';
bb = Xrows \ (x');
xres = Xrows * bb - x';
xres = sum(abs(xres));
cutoff = sqrt(eps(class(xres))) * size(xres,2);
gmeans(xres > cutoff) = NaN;

% Tukey-Kramer
if (isempty(ctype)), ctype = 'tukey-kramer'; end
crit = getcrit(ctype, alpha, stats.dfe, length(gmeans));

ngroups = size(ffdesign,1);
gnames = cell(ngroups,1);
allnames = stats.grpnames;
varnames = stats.varnames;
for j=1:ngroups
    v1 = dim(1);
    vals1 = allnames{v1};
    nm = sprintf('%s=%s',varnames{v1},vals1{ffdesign(j,1)});
    for i=2:size(ffdesign,2)

```

```

v2 = dim(i);
vals2 = allnames{v2};
nm = sprintf('%s,%s=%s',nm,varnames{v2},vals2{ffdesign(j,i)} );
end
gnames{j} = nm;
end

case 'aoctool'
model = stats.model;
if (model==1 || model==3)
error('stats:multcompare:NoMultipleParameters',...
    'No multiple comparisons possible from this aoctool fit.');
end
gnames = stats.gnames;
n = stats.n();
ng = length(n);
df = stats.df;
if (df < 1)
error('stats:multcompare:NotEnoughData',...
    'Cannot do multiple comparison with 0 d.f. for error.');
end

% Estimasi intercept dan covariance
if (isempty(estimate))
if (model == 5)
estimate = 'slope';
else
estimate = 'intercept';
end
end

if (isequal(estimate,'s')), estimate = 'slope'; end
if (isequal(estimate,'i')), estimate = 'intercept'; end
if (isequal(estimate,'p')), estimate = 'pmm'; end
if (~isempty(estimate))
if ~(isequal(estimate,'slope') || isequal(estimate,'pmm')...
    || isequal(estimate,'intercept'))
error('stats:multcompare:BadEstimate',...
    'ESTIMATE must be "slope", "intercept", or "pmm".');
end
end
switch(estimate)
case 'slope'
if (~isfield(stats, 'slopes'))
error('stats:multcompare:BadStats',...
    'No slope estimates available in STATS argument.');
end
gmeans = stats.slopes;

```

```

gcov = stats.slopecov;
mname = 'slopes';
case 'intercept'
if (~isfield(stats, 'intercepts'))
    error('stats:multcompare:BadStats',...
        'No intercept estimates available in STATS argument.');
end
gmeans = stats.intercepts;
gcov = stats.intercov;
mname = 'intercepts';
case 'pmm'
gmeans = stats.pmm;
gcov = stats.pmmcov;
mname = 'population marginal means';
end

if (any(any(isinf(gcov))))
    error('stats:multcompare:InfiniteVariance',...
        'Cannot do comparisons. Some %s have infinite variance.', mname);
end

% Tukey-Kramer critical value
if (isempty(ctype)), ctype = 'tukey-kramer'; end
crit = getcrit(ctype, alpha, df, ng);

case 'kruskalwallis'
gmeans = stats.meanranks(:);
gnames = stats.gnames;
n = stats.n(:);
sumt = stats.sumt;
ng = length(n);
N = sum(n);
mname = 'mean ranks';

if (isempty(ctype)), ctype = 'tukey-kramer'; end
crit = getcrit(ctype, alpha, Inf, ng);

gcov = diag(((N*(N+1)/12) - (sumt/(12*(N-1)))) ./ n);

if (dodisp)
    disp(['Note: Intervals can be used for testing but are not',...
        'simultaneous confidence intervals.']);
end

case 'friedman'
gmeans = stats.meanranks(:);
n = stats.n;
ng = length(gmeans);

```

```

sigma = stats.sigma;
mname = 'mean column ranks';
gnames = strjust(num2str((1:ng)'), 'left');

% Tukey-Kramer
if (isempty(ctype)), ctype = 'tukey-kramer'; end
crit = getcrit(ctype, alpha, Inf, ng);

gcov = ((sigma^2) / n) * eye(ng);

if (dodisp)
    disp(['Note: Intervals can be used for testing but are not ' ...
        'simultaneous confidence intervals.']);
end

otherwise
error('stats:multcompare:BadStats','Bad SOURCE field in STATS structure');
end

[M,MM,hh] = makeM(gmeans, gcov, crit, gnames, mname, dodisp);

comparison = M;
if (nargout>1), means = MM; end
if (nargout>2), h = hh; end

% -----
function crit = getcrit(ctype, alpha, df, ng)

crit = Inf;
[onetype,ctype] = strtok(ctype);

while(~isempty(onetype))
    if (length(onetype) == 1)
        switch onetype
            case 't', onetype = 'tukey-kramer';
            case 'd', onetype = 'dunn-sidak';
            case 'b', onetype = 'bonferroni';
            case 's', onetype = 'scheffe';
            case 'h', onetype = 'tukey-kramer';
            case 'T', onetype = 'lsd';
        end
    end
    if (isequal(onetype, 'hsd')), onetype = 'tukey-kramer'; end

    switch onetype
        case 'tukey-kramer' % or hsd
            crit1 = stdinv(1-alpha, df, ng) / sqrt(2);
    end
end

```

```

ub = getcrit('dunn-sidak', alpha, df, ng);
if (crit1 > ub), crit1 = ub; end

case 'dunn-sidak'
kstar = nchoosek(ng, 2);
alf = 1-(1-alpha).^(1/kstar);
if (isinf(df))
    crit1 = norminv(1-alf/2);
else
    crit1 = tinv(1-alf/2, df);
end

case 'bonferroni'
kstar = nchoosek(ng, 2);
if (isinf(df))
    crit1 = norminv(1 - alpha / (2*kstar));
else
    crit1 = tinv(1 - alpha / (2*kstar), df);
end

case 'lsd'
if (isinf(df))
    crit1 = norminv(1 - alpha / 2);
else
    crit1 = tinv(1 - alpha / 2, df);
end

case 'scheffe'
if (isinf(df))
    tmp = chi2inv(1-alpha, ng-1) / (ng-1);
else
    tmp = finv(1-alpha, ng-1, df);
end
crit1 = sqrt((ng-1) * tmp);

otherwise
error('stats:multcompare:Bad CType',...
    'Unknown CType (critical value type) %s.', ctype);
end

if (~isnan(crit1)), crit = min(crit, crit1); end
[onetype,ctype] = strtok(ctype);
end

% -----
function [M,MM,hh] = makeM(gmeans, gcov, crit, gnames, mname, dodisp)
% Membuat matrix rata-rata dan tampilan

```

```

t = isnan(gmeans);
if any(t)
    gcov(t,:) = 0;
    gcov(:,t) = 0;
end

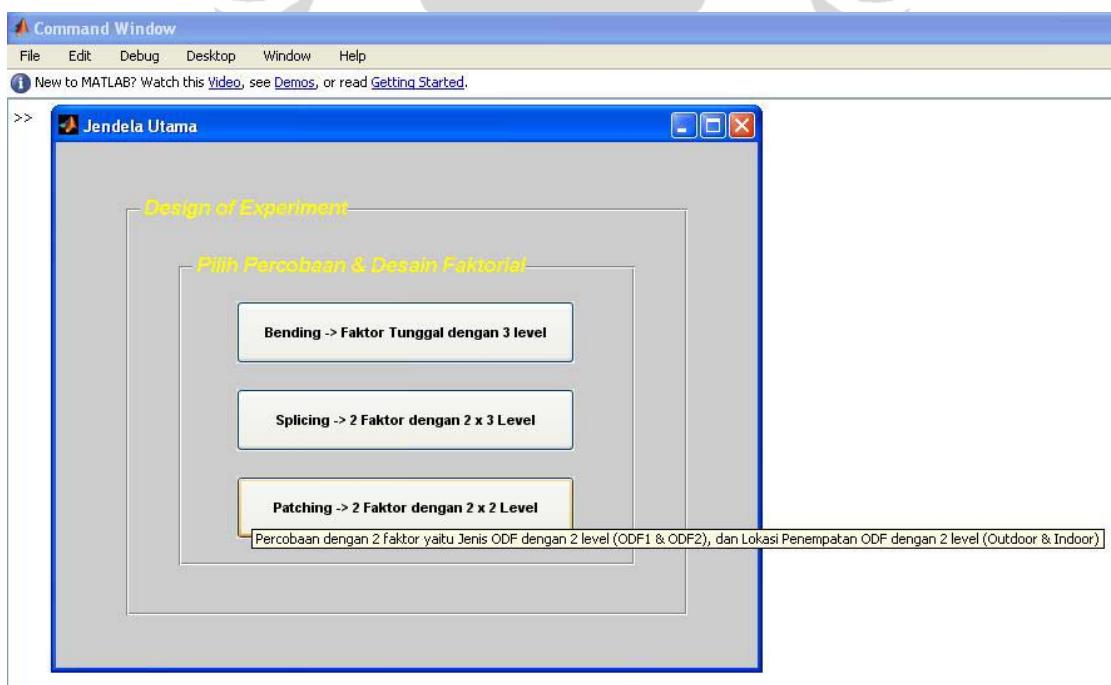
ng = length(gmeans);
MM = zeros(ng,2);
MM(:,1) = gmeans(:,1);
MM(:,2) = sqrt(diag(gcov));
MM(isnan(MM(:,1)),2) = NaN;

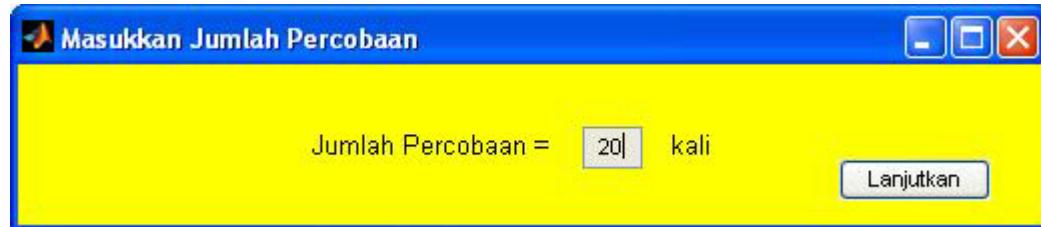
M = nchoosek(1:ng, 2);
M(1,5) = 0;
g1 = M(:,1);
g2 = M(:,2);
M(:,4) = gmeans(g1) - gmeans(g2);
i12 = sub2ind(size(gcov), g1, g2);
gvar = diag(gcov);
d12 = sqrt(gvar(g1) + gvar(g2) - 2 * gcov(i12));
delta = crit * d12;
M(:,3) = M(:,4) - delta;
M(:,5) = M(:,4) + delta;

if (dodisp)
    d = zeros(ng, ng);
    d(i12) = d12;
    sum1 = sum(sum(d));
    d = d + d';
    sum2 = sum(d);
    if (ng > 2)
        w = ((ng-1) * sum2 - sum1) ./ ((ng-1)*(ng-2));
    else
        w = repmat(sum1, 2, 1) / 2;
    end
    halfwidth = crit * w(:,1);
    hh = meansgraph(gmeans, gmeans-halfwidth, gmeans+halfwidth, ...
                    gnames, mname);
    set(hh, 'Name', sprintf('Multiple comparison of %s',mname),'NumberTitle','off');
else
    hh = [];
end

```

Lampiran 2. Tampilan Program Simulasi





Data Percobaan Bending

Kombinasi Level (L)-

	Brand 1	Brand 2	Brand 3
1	0.0370	0.0560	0.1018
2	0.0150	0.0400	0.0854
3	0.0070	0.0150	0.1016
4	0.0600	0.0040	0.1084
5	0.0160	0.0680	0.0652
6	0.0730	0.0690	0.1258
7	0.0490	0.0350	0.0752
8	0.0530	0.0250	0.0985
9	0.0170	0.0160	0.0057
10	0.0400	0.0090	0.0754
11	0.0120	0.1010	0.0512
12	0.0260	0.0150	0.0421
13	0.0690	0.0500	0.0092
14	0.0590	0.0690	0.0026
15	0.0710	0.0780	0.0671

Analisa Data

1. Uji Normalitas

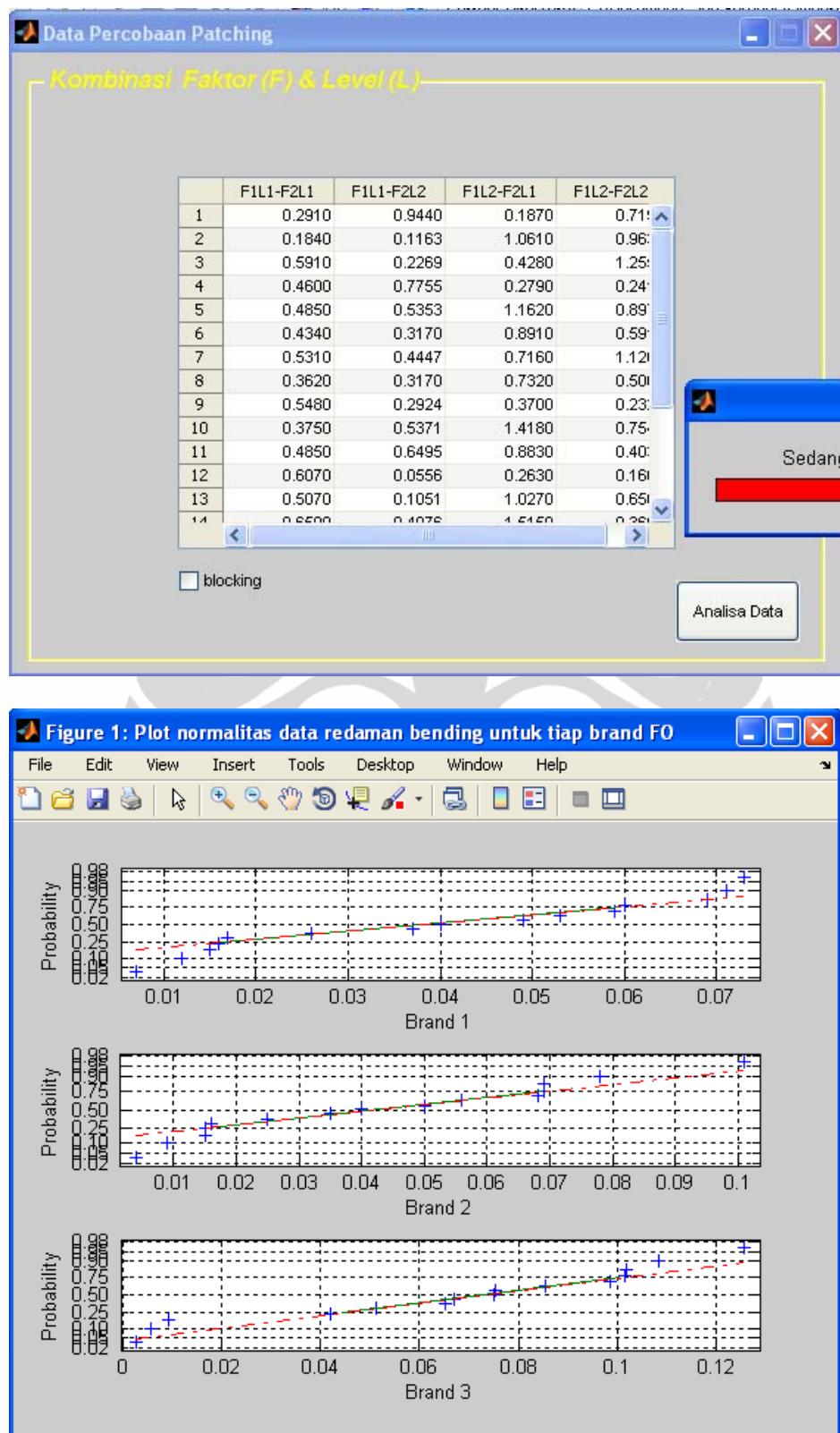
Data Percobaan Splicing

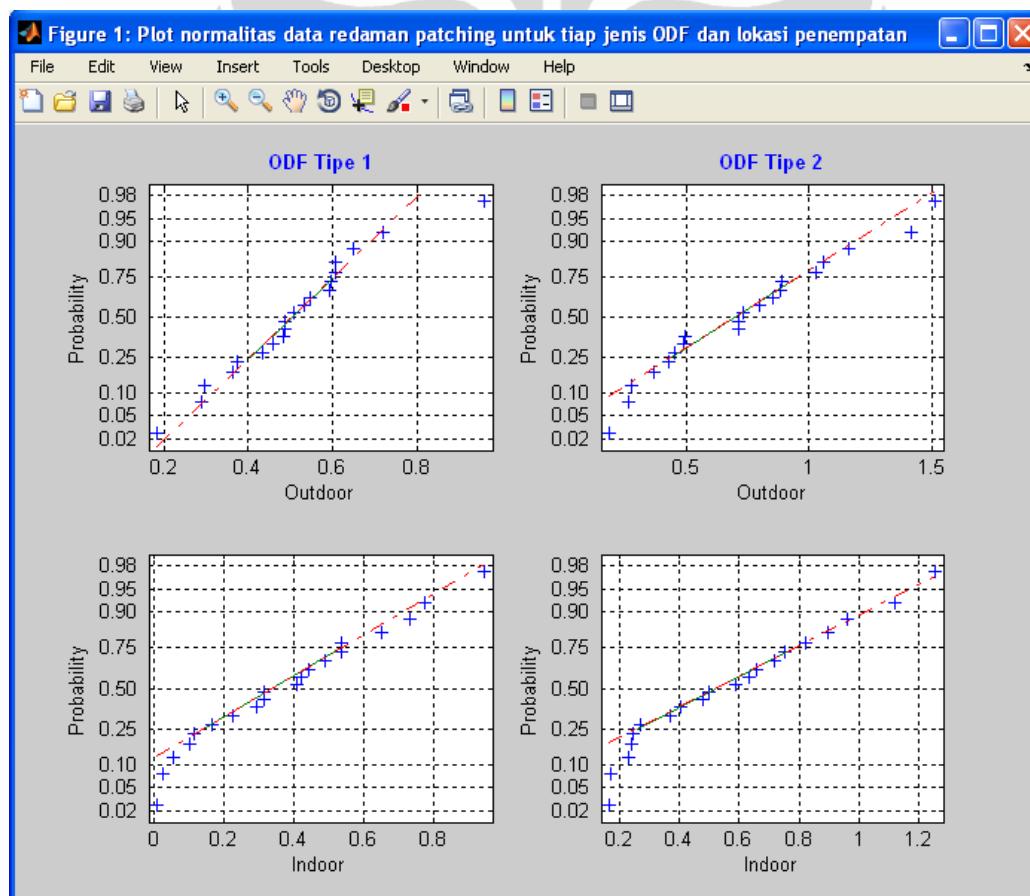
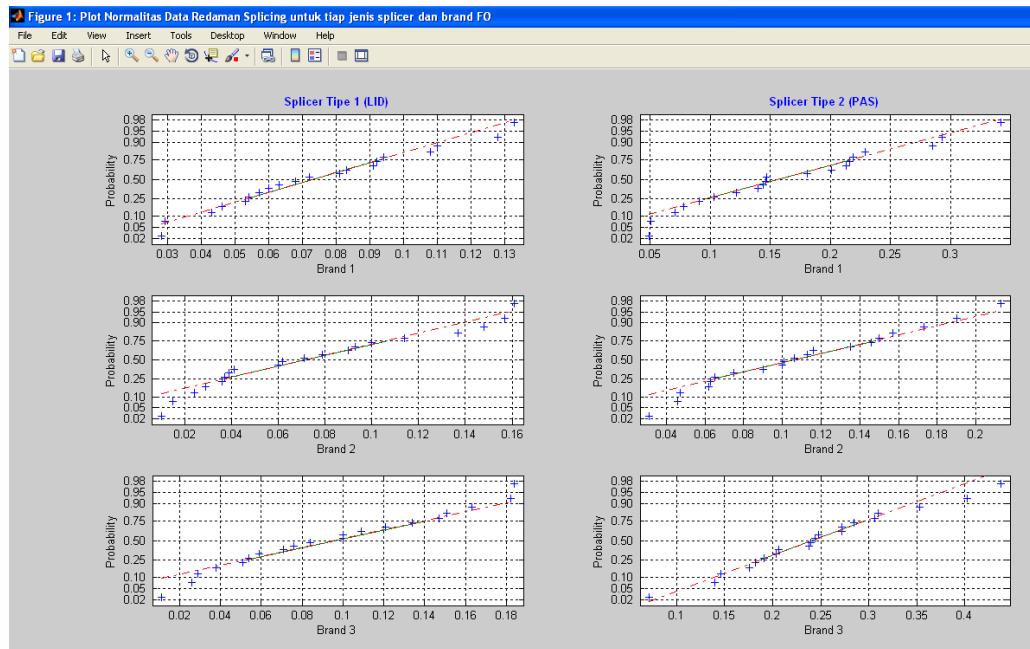
Kombinasi Faktor (F) & Level (L)

	F1L1-F2L1	F1L1-F2L2	F1L1-F2L3	F1L2-F2L1
1	0.0570	0.1610	0.0760	0.21
2	0.0290	0.1480	0.1090	0.28
3	0.0920	0.0100	0.1510	0.12
4	0.0430	0.0790	0.0590	0.20
5	0.0810	0.0930	0.0540	0.14
6	0.0460	0.0390	0.1210	0.34
7	0.0530	0.1370	0.0710	0.29
8	0.0600	0.0620	0.0260	0.14
9	0.0280	0.1140	0.1340	0.07
10	0.0720	0.1570	0.1470	0.21
11	0.0680	0.0290	0.1630	0.14
12	0.0630	0.0150	0.0290	0.18
13	0.1280	0.0370	0.1820	0.09
14	0.1320	0.0710	0.1000	0.24

blocking

Analisa Data





Lampiran 3. Spesifikasi OTU-8000

Specifications	
Base Unit Technical	
Mechanical	
Height	2 RU
Width	19-, 21- (ETSI), or 23-in
Depth	260 mm (ETSI), 300 mm (19- or 23-in)
Power Supply	
DC input	-36 to -60 V
Power consumption	30 W
Environmental	
Operating	-10 to 50°C
Storage	-20 to 60°C
Relay Contacts (Option)	
Humidity	95% without condensing
EMI/ESD	CE Compliant
Interfaces	
1 RJ45 Ethernet 10/100BaseT Port	
1 RJ11 if equipped with PSTN modem	
GSM if equipped with GSM modem	
Storage	
Media	Solid state disk
Optical links (max)	512
Alarms storage	512
OTDR Trace storage	1024
OTDR	
The OTU-8000 can house two field-interchangeable OTDR modules. A wide range of OTDRs are available, ensuring optimum monitoring of all types of fiber optic networks from short-range multimode to long-haul single-mode. The OTU-8000 monitors active fibers using the 1625 nm OTDR module, which allows for factors such as the Raman effect of the optical amplifier.	
Technical	
Distance Unit	km, kft, miles
Group Index Range	1.30000 to 1.70000 in 0.00001 steps
No. of Data points	Up to 128 000 data points
Distance Measurement	Automatic or dual cursor
Display span	From 2.6 m up 380 km
Display resolution	1 cm
Cursor resolution	From 1 cm
Sampling resolution	From 4 cm
Accuracy	$\pm 1 \text{ m} \pm \text{sampling resolution}$ $\pm 1 \cdot 10^{-5} \times \text{distance}$ (excluding group index uncertainties)
Attenuation Measurement	
Automatic, manual, 2-points, 5-points and LSA	
Display span	1.25 to 55 dB
Display resolution	0.001 dB
Cursor resolution	From 0.001 dB
Accuracy	$\pm 0.05 \text{ dB} \pm 0.05 \text{ dB/PI}$
Reflectance/ORL Measurements	
Automatic or manual	
Display resolution	0.01 dB
Threshold	-11 to -99 dB in 1 dB step
Optical Switch	
The OTU-8000 can house a field interchangeable optical switch module with up to 24 ports. If higher ports count is required, 24 ports can be extended to more than 1000 ports by adding chassis of 64 ports each. An OTU-8000 with no OTDR modules fitted forms the base of the Remote Optical Switch controlled by TCP/IP.	
Upgrade the Remote Optical Switch base by adding an OTDR module to become a complete OTU-8000 at any time.	
General	
No. of Ports	2, 4, 8, 12, 16, 24, 36, 48, or 64 with 1 or 2 commons, more than 1000 ports with 1 common
Insertion Loss(excluding connectors)	0.6 dB up to 64 ports; 1.2 dB for higher port counts
Back-Reflection	-60 dB (Single mode)
Repeatability	$\pm 0.01 \text{ dB}$
Wavelength Range	1310, 1550 and 1625 nm
Lifetime	10^7 cycles
Housing	Up to 24 ports: Included in the OTU-8000 For higher port counts: external 4 RU racks

The main specifications of OTDR modules are available on the consolidated OTDR datasheet.



Lampiran 4. Spesifikasi ITU-T G.652 D

Table 4/G.652 – G.652.D attributes

Fibre attributes		
Attribute	Detail	Value
Mode field diameter	Wavelength	1310 nm
	Range of nominal values	8.6-9.5 μm
	Tolerance	$\pm 0.6 \mu\text{m}$
Cladding diameter	Nominal	125.0 μm
	Tolerance	$\pm 1 \mu\text{m}$
Core concentricity error	Maximum	0.6 μm
Cladding noncircularity	Maximum	1.0%
Cable cut-off wavelength	Maximum	1260 nm
Macrobend loss	Radius	30 mm
	Number of turns	100
	Maximum at 1625 nm	0.1 dB
Proof stress	Minimum	0.69 GPa
Chromatic dispersion coefficient	$\lambda_{0\min}$	1300 nm
	$\lambda_{0\max}$	1324 nm
	$S_{0\max}$	0.092 ps/nm ² × km
Cable attributes		
Attribute	Detail	Value
Attenuation coefficient	Maximum from 1310 nm to 1625 nm (Note 2)	0.4 dB/km
	Maximum at 1383 nm ± 3 nm	(Note 3)
	Maximum at 1550 nm	0.3 dB/km
PMD coefficient	M	20 cables
	Q	0.01%
	Maximum PMD _Q	0.20 ps/ $\sqrt{\text{km}}$
<p>NOTE 1 – According to 6.2, a maximum PMD_Q value on uncabled fibre is specified in order to support the primary requirement on cable PMD_Q.</p> <p>NOTE 2 – This wavelength region can be extended to 1260 nm by adding 0.07 dB/km induced Rayleigh scattering loss to the attenuation value at 1310 nm. In this case, the cable cut-off wavelength should not exceed 1250 nm.</p> <p>NOTE 3 – The sampled attenuation average at this wavelength shall be less than or equal to the maximum value specified for the range, 1310 nm to 1625 nm, after hydrogen ageing according to IEC 60793-2-50 regarding the B1.3 fibre category.</p>		

Lampiran 5. Tabel Konversi Daya Redaman

Redaman (dB)	Persentasi Daya yang Diterima di Penerima	Persentasi Daya yang Hilang	Keterangan
1	79%	21%	---
2	63%	37%	---
3	50%	50%	1/2 Daya Pengirim
4	40%	60%	---
5	32%	68%	---
6	25%	75%	1/4 Daya Pengirim
7	20%	80%	1/5 Daya Pengirim
8	16%	84%	1/6 Daya Pengirim
9	12%	88%	1/8 Daya Pengirim
10	10%	90%	1/10 Daya Pengirim
11	8%	92%	1/12 Daya Pengirim
12	6.30%	93.70%	1/16 Daya Pengirim
13	5%	95%	1/20 Daya Pengirim
14	4%	96%	1/25 Daya Pengirim
15	3.20%	96.80%	1/30 Daya Pengirim
16	2.50%	97.50%	1/40 Daya Pengirim
17	2%	98%	1/50 Daya Pengirim
18	1.60%	98.40%	1/60 Daya Pengirim
19	1.30%	98.70%	1/80 Daya Pengirim
20	1%	99%	1/100 Daya Pengirim
25	0.30%	99.70%	1/300 Daya Pengirim
30	0.10%	99.90%	1/1000 Daya Pengirim
40	0.01%	99.99%	1/10,000 Daya Pengirim
50	0.00%	100.00%	1/100,000 Daya Pengirim