

Auswertung mit Python

August 24, 2025

1 Auswertung 27.05.2025

Auswertung mit Vergleich von TBBB TCBC und LNO

1.1 Präambel

```
[181]: import numpy as np
import matplotlib.pyplot as plt
from matplotlib import cm
from scipy.optimize import curve_fit

import glob
import re

def dat2array(file_path):
    data = []
    with open(file_path, 'r', encoding='utf-8') as file:
        for i, line in enumerate(file):
            line = line.replace(',', '.').replace('\t', ';').replace('\n', '')
            if i >= 39: # Skip header lines
                data.append(line.split(';'))

    freqs = [float(row[0]) for row in data]
    intensities = [float(row[1]) for row in data]
    return freqs, intensities

def extract_info(filename):
    match = re.search(r'0(\d+)_(\d+)K.asc', filename)
    if match:
        return int(match.group(1)), int(match.group(2))
    return None, None

def gaussian(x, a, mean, sigma):
    return a * np.exp(-(x - mean) ** 2 / (2 * sigma ** 2))
```

```
[182]: # Hilfsfunktion: FWHM aus diskreten Daten via linearer Interpolation bestimmen
def _compute_fwhm_from_discrete(x, y, ymax=None):
    if x is None or y is None or len(x) == 0 or len(y) == 0:
```

```

        return None
    if ymax is None:
        ymax = max(y)
    half = ymax / 2.0
    idx_max = int(np.argmax(y))
    # Linke Halbwertsstelle
    left = None
    for i in range(idx_max, 0, -1):
        y1, y2 = y[i-1], y[i]
        if (y1 - half) * (y2 - half) <= 0: # Vorzeichenwechsel bzw. Schnitt
            x1, x2 = x[i-1], x[i]
            if y2 != y1:
                left = x1 + (half - y1) * (x2 - x1) / (y2 - y1)
            else:
                left = x[i]
            break
    # Rechte Halbwertsstelle
    right = None
    for i in range(idx_max, len(y) - 1):
        y1, y2 = y[i], y[i+1]
        if (y1 - half) * (y2 - half) <= 0:
            x1, x2 = x[i], x[i+1]
            if y2 != y1:
                right = x1 + (half - y1) * (x2 - x1) / (y2 - y1)
            else:
                right = x[i]
            break
    if left is None or right is None:
        return None
    return abs(right - left)

```

```

[183]: def create_shg_plot(middle_value, difference, filepath, supitle,
    ↪ filename_prefix, reference=False, compare=True, power="333pW",
    ↪ slit_width="210µm", normalisationfactor=131942273.14285715,
    ↪ refmeas="LiNbO -SHG"):
    x_range = (middle_value - difference, middle_value + difference)
    file_list = glob.glob(filepath)

    sorted_files = []
    for f in file_list:
        meas_no, temperature = extract_info(f)
        if meas_no is not None:
            sorted_files.append((f, meas_no, temperature))

    sorted_files.sort(key=lambda x: x[1])

    extended_files = []

```

```

heating = True
for f, meas_no, temperature in sorted_files:
    if heating and temperature > 435:
        heating = False
    extended_files.append((f, meas_no, temperature, heating))

temperatures = [x[2] for x in sorted_files]
if len(temperatures) == 0:
    print("Keine Dateien gefunden, die dem Pfad entsprechen.")
    return
norm = plt.Normalize(min(temperatures), max(temperatures))
colors = cm.plasma(norm(temperatures))

# Bestimme zuerst das absolute Maximum aus den Rohdaten im gezeigten Fenster
# (vor jeglicher Referenz-/Normalisierungs- bzw. Skalierungsanpassung)
absolute_max_raw = None
for f, meas_no, temperature, heating in extended_files:
    freqs_tmp, ints_tmp = dat2array(f)
    # Rohdaten verwenden - keine Referenz oder Normalisierung anwenden
    vals = [i for fr, i in zip(freqs_tmp, ints_tmp) if x_range[0] <= fr <=
↪x_range[1]]
    if vals:
        local_max = max(vals)
        if absolute_max_raw is None or local_max > absolute_max_raw:
            absolute_max_raw = local_max
if absolute_max_raw is None:
    absolute_max_raw = 0.0
print(f"Absolute maximum in raw data within x_range {x_range}:
↪{absolute_max_raw}")

# Wenn compare=False: zähle zusätzlich das globale Maximum im gezeigten
↪Fenster (kann andere Skalierung nutzen)
global_max = 0.0
if not compare:
    for f, meas_no, temperature, heating in extended_files:
        freqs_tmp, ints_tmp = dat2array(f)
        if reference:
            ints_tmp = [val * 32.88 * (10/7) for val in ints_tmp]
            # Filter auf das gezeigte Fenster
            vals = [i for fr, i in zip(freqs_tmp, ints_tmp) if x_range[0] <= fr
↪<= x_range[1]]
            if vals:
                local_max = max(vals)
                if local_max > global_max:
                    global_max = local_max
if global_max <= 0:
    global_max = 1.0

```

```

fig, (ax0, ax1) = plt.subplots(1, 2, figsize=(15, 5))
peak_list = []
temps_for_line = []
peaks_for_line = []

# Sammle Peak-Parameter je Messung für spätere Zusammenfassung
peak_metrics = [] # Einträge: dict(height, area, mean, fwhm, temperature,
↳ fit_success)

for idx, (f, meas_no, temperature, heating) in enumerate(extended_files):
    freqs, intensities = dat2array(f)
    # optionale Referenzskalierung (z.B. ND2 und Zeitkorrektur)
    if reference:
        intensities = [val * 32.88 * (10/7) for val in intensities]

    # falls Vergleichsmodus an, normalisiere wie vorher relativ zu LND
    ↳ (Konstante bleibt)
    if compare:
        intensities = [val / normalisationfactor for val in intensities]

    # Filtere nur das gezeigte Fenster (wichtig!)
    filtered_data = [(fr, i) for fr, i in zip(freqs, intensities) if
    ↳ x_range[0] <= fr <= x_range[1]]
    filtered_freqs, filtered_intensities = zip(*filtered_data) if
    ↳ filtered_data else ([], [])

    if not filtered_data:
        continue

    # falls compare=False -> skaliere alle Messdaten relativ zum globalen
    ↳ Maximum im Fenster
    if not compare:
        if global_max <= 0:
            scale = 1.0
        else:
            scale = global_max
        filtered_intensities = [v / scale for v in filtered_intensities]
    else:
        filtered_intensities = list(filtered_intensities)

    filtered_freqs = list(filtered_freqs)

    ax0.plot(filtered_freqs, filtered_intensities, color='lightgray') #
    ↳ Plot in Hellgrau

```

```

    try:
        popt, _ = curve_fit(gaussian, filtered_freqs, filtered_intensities,
                             p0=[max(filtered_intensities), np.
→mean(filtered_freqs), 10])
        fitted_curve = gaussian(np.array(filtered_freqs), *popt)
        ax0.plot(filtered_freqs, fitted_curve, color=colors[idx])
        area = np.trapezoid(fitted_curve, filtered_freqs)
        # --- Hier erfolgt die x-Achsen-Anpassung ---
        temp_for_plot = temperature # Standardwert
        if not heating:
            temp_for_plot = 880 - temperature
        ax1.scatter(temperature, area, color='black', marker='o')

        # Peak-Parameter aus Fit
        a_fit, mean_fit, sigma_fit = float(popt[0]), float(popt[1]),
→abs(float(popt[2]))
        fwhm_fit = 2.0 * np.sqrt(2.0 * np.log(2.0)) * sigma_fit
        peak_metrics.append({
            'height': a_fit,
            'area': float(area),
            'mean': mean_fit,
            'fwhm': float(fwhm_fit),
            'temperature': temperature,
            'fit_success': True,
        })
    except RuntimeError:
        ax0.plot(filtered_freqs, filtered_intensities, color=colors[idx])
        area = np.trapezoid(filtered_intensities, filtered_freqs)
        ax1.scatter(temperature, area, color='black', marker='x') #
→Original auskommentieren

        # Peak-Parameter ohne Fit (robuste Schätzung)
        if filtered_intensities and filtered_freqs:
            height_nf = float(max(filtered_intensities))
            mean_nf = float(filtered_freqs[int(np.
→argmax(filtered_intensities))])

            # Lokale FWHM-Schätzung per Interpolation
            def _compute_fwhm_from_discrete_local(x, y, ymax=None):
                if x is None or y is None or len(x) == 0 or len(y) == 0:
                    return None
                if ymax is None:
                    ymax = max(y)
                half = ymax / 2.0
                idx_max = int(np.argmax(y))
                left = None
                for i in range(idx_max, 0, -1):
                    y1, y2 = y[i-1], y[i]

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        if (y1 - half) * (y2 - half) <= 0:
            x1, x2 = x[i-1], x[i]
            if y2 != y1:
                left = x1 + (half - y1) * (x2 - x1) / (y2 - y1)
            else:
                left = x[i]
            break
    right = None
    for i in range(idx_max, len(y) - 1):
        y1, y2 = y[i], y[i+1]
        if (y1 - half) * (y2 - half) <= 0:
            x1, x2 = x[i], x[i+1]
            if y2 != y1:
                right = x1 + (half - y1) * (x2 - x1) / (y2 - y1)
            else:
                right = x[i]
            break
    if left is None or right is None:
        return None
    return abs(right - left)
    fwhm_nf = _compute_fwhm_from_discrete_local(list(filtered_freqs),
    list(filtered_intensities), ymax=height_nf)
    peak_metrics.append({
        'height': height_nf,
        'area': float(area),
        'mean': mean_nf,
        'fwhm': float(fwhm_nf) if fwhm_nf is not None else None,
        'temperature': temperature,
        'fit_success': False,
    })

    peak_list.append(area)
    temps_for_line.append(temperature)
    peaks_for_line.append(area)

sm = plt.cm.ScalarMappable(cmap=cm.plasma, norm=norm)
sm.set_array([])
cbar = plt.colorbar(sm, ax=ax0, orientation='vertical', pad=0.02)
cbar.set_label('Temperature in K')
cbar.set_ticks([min(temperatures), max(temperatures)])
cbar.set_ticklabels([f"{min(temperatures)} K", f"{max(temperatures)} K"])

# Finde Index, an dem heating von True auf False wechselt
heating_flags = [ht for (_, _, _, ht) in extended_files]
if True in heating_flags and False in heating_flags:
    split_idx = heating_flags.index(False)

```

```

ax1.plot(temps_for_line[:split_idx+1], peaks_for_line[:split_idx+1],
         color='red', linewidth=1.2, alpha=0.5, zorder=0)
ax1.plot(temps_for_line[split_idx:], peaks_for_line[split_idx:],
         color='blue', linewidth=1.2, alpha=0.5, zorder=0)
else:
    color = 'red' if all(heating_flags) else 'blue'
    ax1.plot(temps_for_line, peaks_for_line, color=color, linewidth=2,
    ↪alpha=0.7, zorder=0)

# set x- and y-axis limits and labels
ax0.set_xlim(x_range)
ax0.set_ylim(bottom=0)
ax0.set_xlabel('Wavelength in nm')

# Achsenbeschriftung abhängig vom compare-Flag
if compare:
    ax0.set_ylabel(f'rel. Int. to {refmeas}')
    ax1.set_ylabel(f'Area of Gaussian Fit rel. to {refmeas}')
else:
    ax0.set_ylabel('Intensity in arb. u.')
    ax1.set_ylabel('Area of Gaussian Fit in arb. u.')

fig.suptitle(suptitle, fontsize=16)
fig.text(0.5, 0.914, f"Measurement parameters: Power: {power}, Slit_
↪width={slit_width}", ha="center")

# Zusammenfassung unterhalb der Diagramme anzeigen
if len(peak_metrics) > 0:
    # Peak für die Zusammenfassung anhand der größten Fläche wählen (nur im
    ↪gezeigten Fenster berechnet)
    best = max(peak_metrics, key=lambda d: d.get('area', float('-inf')))
    lam0 = best.get('mean', None)
    fwhm = best.get('fwhm', None)
    hoehe = best.get('height', None)
    flaeche = best.get('area', None)
    unit = 'arb. u.' if (reference or not compare) else 'rel.'
    parts = []
    if lam0 is not None:
        parts.append(f"Zentralwellenlänge: {lam0:.2f} nm")
    if fwhm is not None:
        parts.append(f"FWHM: {fwhm:.2f} nm")
    # Bei compare=False sollen Höhe und Fläche im Info-Kasten weggelassen
    ↪werden
    if compare:
        if hoehe is not None:
            parts.append(f"Höhe: {hoehe:.3g} {unit}")
        if flaeche is not None:

```

```

        parts.append(f"Fläche: {flaeche:.3g} {unit}·nm")

    info_text = "    |    ".join(parts) if parts else ""
    if info_text:
        fig.subplots_adjust(bottom=0.18)
        fig.text(0.5, 0.01, info_text, ha='center', va='bottom',
                fontsize=10,
                bbox=dict(boxstyle='round', facecolor='white', alpha=0.9,
                        edgecolor='lightgray'))

plt.show()

fig.savefig(f'./{filename_prefix}.png', dpi=600, bbox_inches='tight')
fig.savefig(f'./{filename_prefix}.pdf', dpi=600, bbox_inches='tight')

```

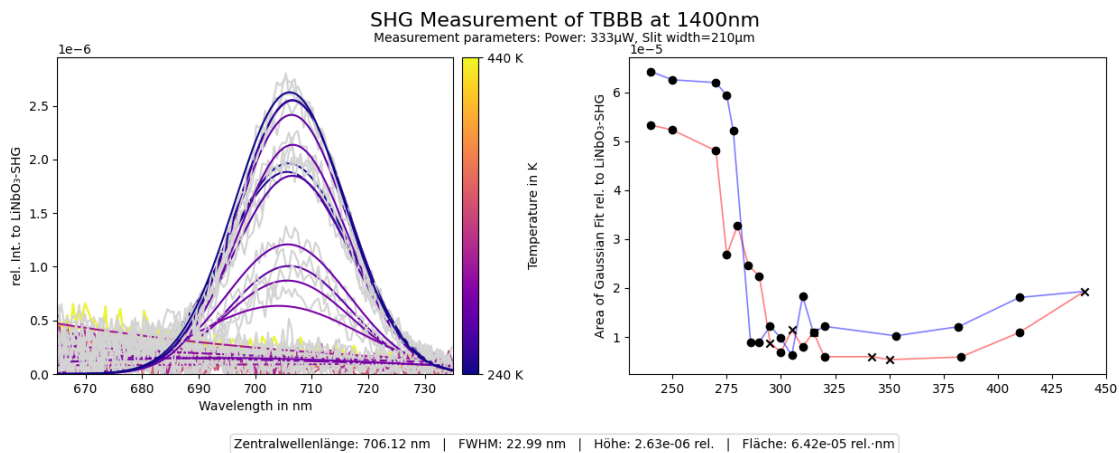
1.2 TBBB

```

[184]: create_shg_plot(
        middle_value=700,
        difference=35,
        filepath='2025-05-22 TBBB/1400nm/0.333mW_Spalt210um/*.asc',
        subtitle='SHG Measurement of TBBB at 1400nm',
        filename_prefix='TBBB_SHG'
    )

```

Absolute maximum in raw data within x_range (665, 735): 370.0



```

[185]: create_shg_plot(
        middle_value=467,
        difference=35,
        filepath='2025-05-22 TBBB/1400nm/0.333mW_Spalt210um/*.asc',

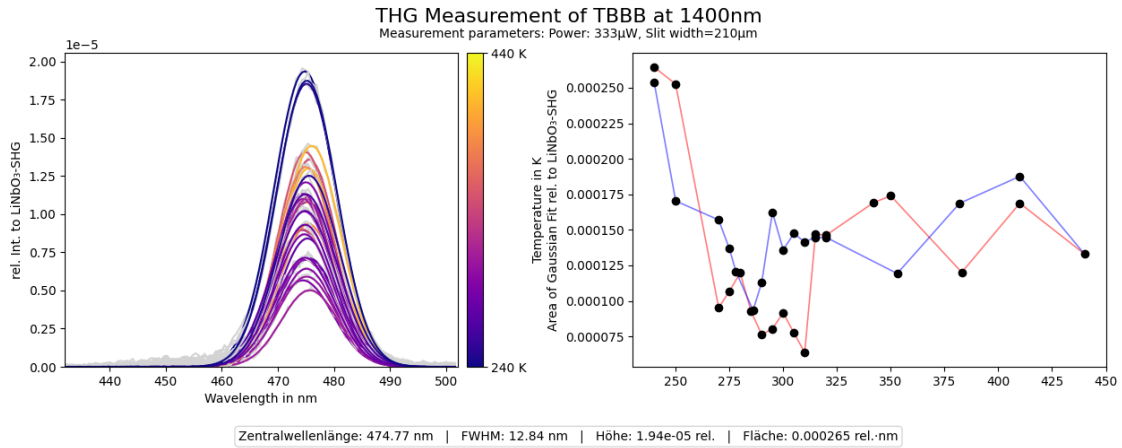
```



```
suptitle='THG Measurement of TBBB at 1400nm',
filename_prefix='TBBB_THG'
```

```
)
```

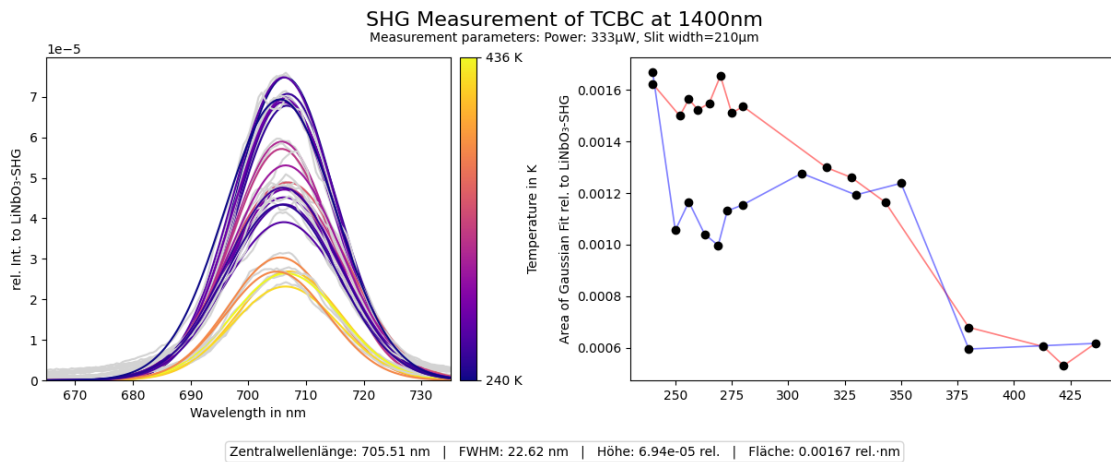
Absolute maximum in raw data within x_range (432, 502): 2580.0



1.3 TCBC

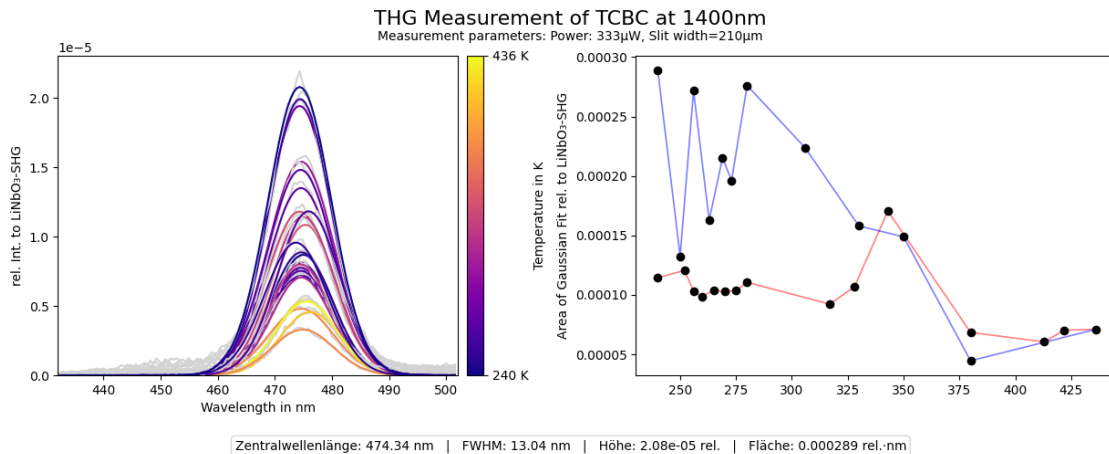
```
[186]: create_shg_plot(
    middle_value=700,
    difference=35,
    filepath='2025-05-23 TCBC/1400nm/0.333mW_Spalt210um/*.asc',
    suptitle='SHG Measurement of TCBC at 1400nm',
    filename_prefix='TCBC_SHG'
)
```

Absolute maximum in raw data within x_range (665, 735): 10023.0



```
[187]: create_shg_plot(
    middle_value=467,
    difference=35,
    filepath='2025-05-23 TCBC/1400nm/0.333mW_Spalt210um/*.asc',
    subtitle='THG Measurement of TCBC at 1400nm',
    filename_prefix='TCBC_THG'
)
```

Absolute maximum in raw data within x_range (432, 502): 2894.0



1.4 LNO

Vergleichsmessung

```
[188]: # Definiere den Wellenlängenbereich wie bei LNO SHG
x_range = (700 - 35, 700 + 35) # anpassen falls verändert!

# Lade die Vergleichsmessungen "mit" und "ohne" ND2.0:
files_ohne = glob.glob('2025-05-23 LNO/1400nm/0.333mW_Spalt210um/
↳Vergleichsmessung ND2.0/ohne*.asc')
files_mit = glob.glob('2025-05-23 LNO/1400nm/0.333mW_Spalt210um/
↳Vergleichsmessung ND2.0/mit*.asc')

# Wähle jeweils die erste Datei (ggf. anpassen, falls mehrere vorhanden)
file_ohne = files_ohne[0]
file_mit = files_mit[0]

# Lade die Daten
freqs_ohne, intensities_ohne = dat2array(file_ohne)
freqs_mit, intensities_mit = dat2array(file_mit)
```

```

# Filtere den Bereich wie bei x_range
filtered_data_ohne = [(fr, i) for fr, i in zip(freqs_ohne, intensities_ohne) if
    ↳x_range[0] <= fr <= x_range[1]]
filtered_data_mit = [(fr, i) for fr, i in zip(freqs_mit, intensities_mit) if
    ↳x_range[0] <= fr <= x_range[1]]

filtered_freqs_ohne, filtered_intensities_ohne = zip(*filtered_data_ohne) if
    ↳filtered_data_ohne else ([], [])
filtered_freqs_mit, filtered_intensities_mit = zip(*filtered_data_mit) if
    ↳filtered_data_mit else ([], [])

fig, ax0 = plt.subplots(figsize=(8, 5))
ax0.plot(filtered_freqs_ohne, filtered_intensities_ohne, label='without ND2.0')
ax0.plot(filtered_freqs_mit, filtered_intensities_mit, label='with ND2.0')
ax0.set_title('Comparison: without ND2.0 vs. with ND2.0')
ax0.set_xlabel('Wavelength in nm')
ax0.set_ylabel('Intensity in arb.u.')
ax0.set_yscale('log')
ax0.legend()

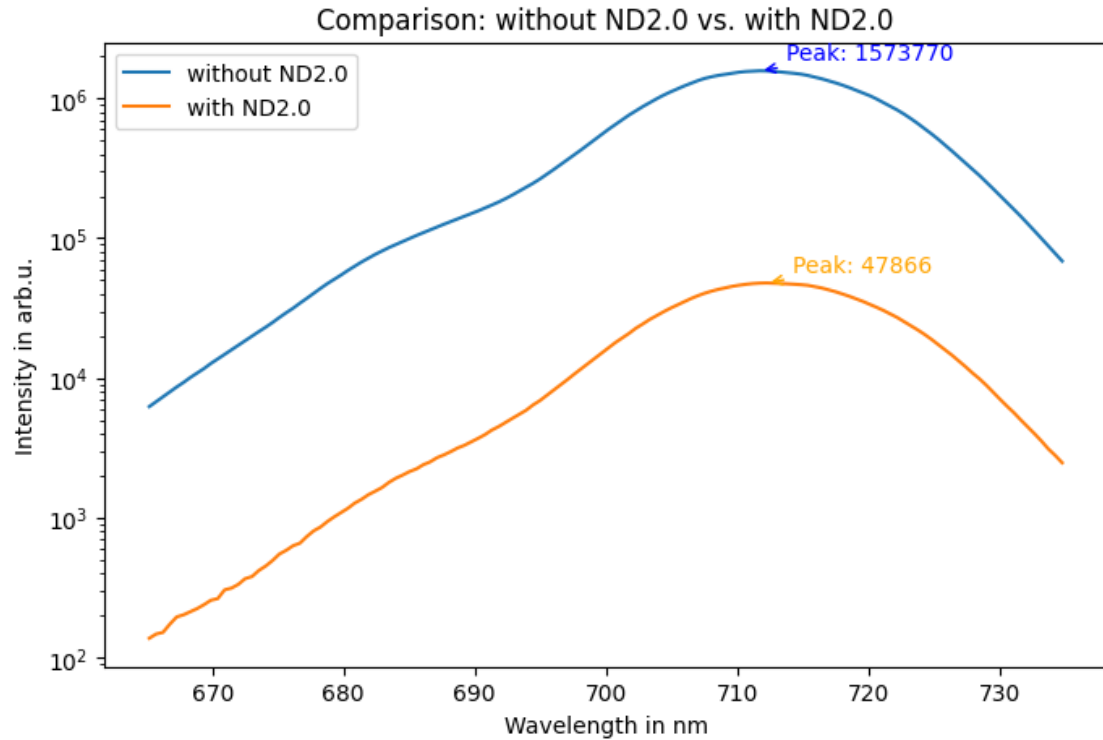
peak_ohne = max(filtered_intensities_ohne)
peak_mit = max(filtered_intensities_mit)
peak_ohne_x = filtered_freqs_ohne[filtered_intensities_ohne.index(peak_ohne)]
peak_mit_x = filtered_freqs_mit[filtered_intensities_mit.index(peak_mit)]

ax0.annotate(f'Peak: {peak_ohne:.0f}', xy=(peak_ohne_x, peak_ohne),
    ↳xytext=(peak_ohne_x+2, peak_ohne*1.2),
    ↳arrowprops=dict(arrowstyle='->', color='blue'), color='blue',
    ↳fontSize=10)
ax0.annotate(f'Peak: {peak_mit:.0f}', xy=(peak_mit_x, peak_mit),
    ↳xytext=(peak_mit_x+2, peak_mit*1.2),
    ↳arrowprops=dict(arrowstyle='->', color='orange'), color='orange',
    ↳fontSize=10)

plt.show()

# Berechne den Skalierungsfaktor, um die Messung mit ND2.0 auf die ohne ND2.0
    ↳zu bringen
skalierungsfaktor = peak_ohne / peak_mit
print(f"Skalierungsfaktor (ohne ND2.0 / mit ND2.0): {skalierungsfaktor:.2f}")

```

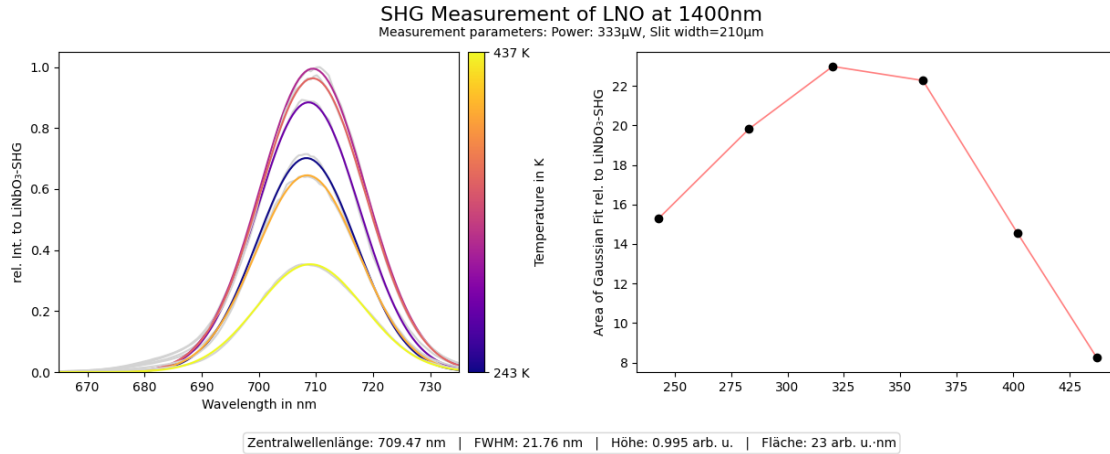


Skalierungsfaktor (ohne ND2.0 / mit ND2.0): 32.88

1.4.1 SHG

```
[189]: create_shg_plot(
    middle_value=700,
    difference=35,
    filepath='2025-05-23 LNO/1400nm/0.333mW_Spalt210um/*.asc',
    subtitle='SHG Measurement of LNO at 1400nm',
    filename_prefix='LNO_SHG',
    reference=True
)
```

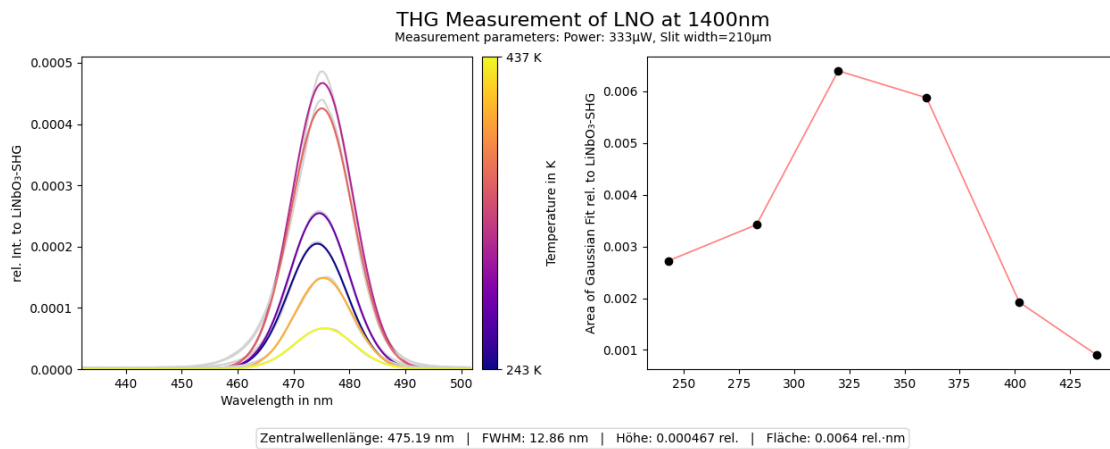
Absolute maximum in raw data within x_range (665, 735): 2808990.0



1.4.2 THG

```
[190]: create_shg_plot(
    middle_value=467,
    difference=35,
    filepath='2025-05-23 LNO/1400nm/0.333mW_Spalt210um/*.asc',
    subtitle='THG Measurement of LNO at 1400nm',
    filename_prefix='LNO_THG'
)
```

Absolute maximum in raw data within x_range (432, 502): 64109.0

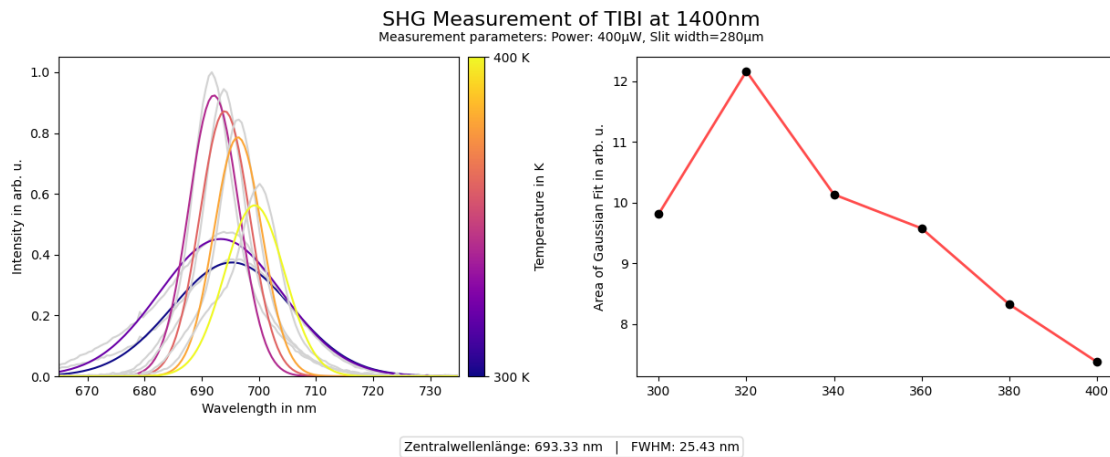


1.5 TIBI

1.5.1 SHG

```
[191]: create_shg_plot(  
    middle_value=700,  
    difference=35,  
    filepath='TIBI_400uW_1400nm/*.asc',  
    suptitle='SHG Measurement of TIBI at 1400nm',  
    filename_prefix='TIBI_SHG',  
    power="400µW",  
    slit_width="280µm",  
    compare=False,  
    reference=False  
)
```

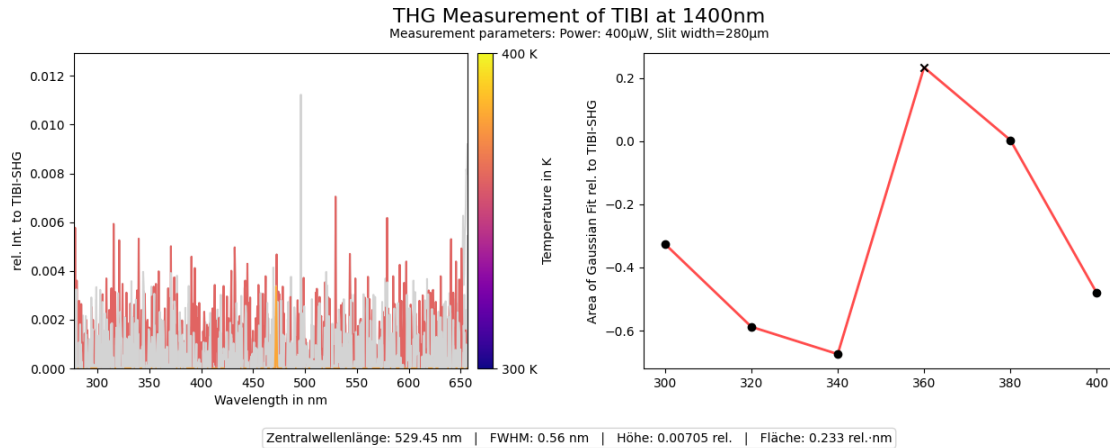
Absolute maximum in raw data within x_range (665, 735): 44702.0



1.5.2 THG

```
[192]: create_shg_plot(  
    middle_value=467,  
    difference=190,  
    filepath='TIBI_400uW_1400nm/*.asc',  
    suptitle='THG Measurement of TIBI at 1400nm',  
    filename_prefix='TIBI_THG',  
    power="400µW",  
    slit_width="280µm",  
    compare=True,  
    normalisationfactor=44702.0,  
    refmeas="TIBI-SHG"  
)
```

Absolute maximum in raw data within x_range (277, 657): 502.0

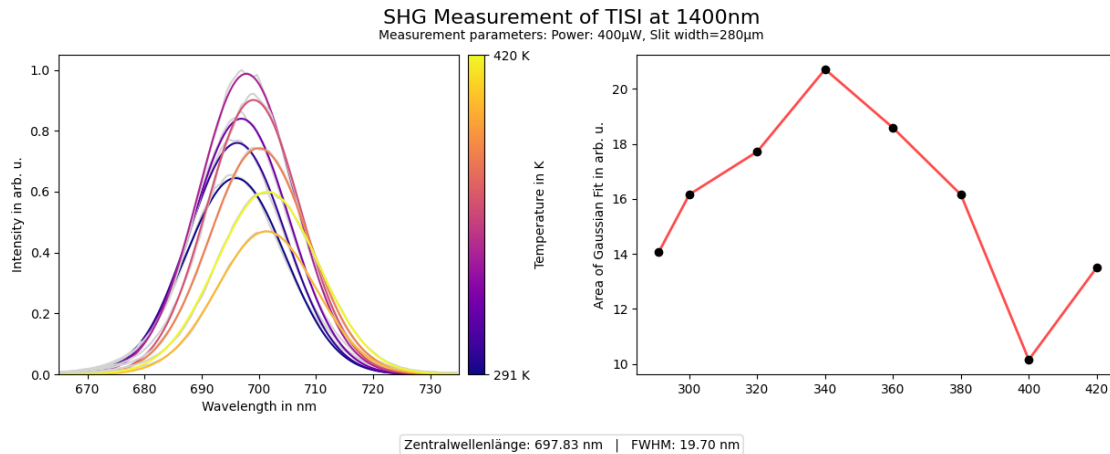


1.6 TISI

1.6.1 SHG

```
[193]: create_shg_plot(
    middle_value=700,
    difference=35,
    filepath='TISI_400uW_1400nm/*.asc',
    suptitle='SHG Measurement of TISI at 1400nm',
    filename_prefix='TISI_SHG',
    power="400µW",
    slit_width="280µm",
    compare=False,
    reference=False
)
```

Absolute maximum in raw data within x_range (665, 735): 162541.0



1.6.2 THG

```
[194]: create_shg_plot(  
    middle_value=467,  
    difference=190,  
    filepath='TISI_400uW_1400nm/*.asc',  
    subtitle='THG Measurement of TISI at 1400nm',  
    filename_prefix='TISI_THG',  
    power="400μW",  
    slit_width="280μm",  
    compare=True,  
    normalisationfactor=162541.0,  
    refmeas="TISI-SHG"  
)
```

Absolute maximum in raw data within x_range (277, 657): 699.0

